SEQ ID	-		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
i	sequence		03/340,21/	sequence	or pepade sequence	deterior, a possible indeteoride insertion,
24216	54584	A	24354	1	1201	MQPALLIVLNKVRQEARRRAR
						ACRIRRASGIVQFGKVRRQLAA
						VELRQRQTPERFVLLRGTRQQA
						TGQRIVKTEQRVIIITQRSFRRPG
						EGRRINDHFRFPAHLPSIRPIGK
						HQTPFGISGHHFNFFTVAIGDDI
						AQLDRIGANQSVLKGFNKFVV
				1		QYATDSMTSQGKGLSQGSGVA
		l				FDNEKFAYNI/SFNGHMLRILDH
		l				G\INWDNDNGTKWWTVGIRPM
						YKWTPIMSTVMEIGYDNVESQ
		1	i			RTGDKNNQYKITLAQQWQAG
ļ		1				DSIWSRPAIRVFATYAKWDEK
						WRYDYTGNADNNANFGKAVP
						ADFNGGSFG\RGDSDEWTFGAQ
						MEIWGLSRASGTLVTRAWRYK
					}	TTPTRHCLLMNHYAGEAAMR
						WKNYLKSTQKVQLQNFANWP
				:	İ	KYSLSTLFHFPFSKSC
24217	54585	Α	24355	188	406	FIRSFDPSFVPSVLDRFFVLSTVF
		1				HSSSIFLVVIDRPSCVGEPSLF*V
						GGSCPTNSLYRCSGLAWYLMR
						SMD
24218	54586	Α	24356	1	1798	
24219	54587	В	24357	1465	7841	
24220	54588	Α	24358	174	3762	TLNDAMRREKCLARINRTAKS
		1				GVVMPKLVTWMNNQRVGELT
						KLANGAHTFKYAPEWLASRYA
		l				RPLSLSLPLQRGNITSDAVFNFF
						DNLLPDRTIVRDRIVKRYHAKS
						RQPFDLLSEIGRDSVGAVTLIPE
						DETQHHYESLTITDEGEFVVFV
		ŀ				GPSGCGKSTLLRMIAGLETITTG
						DLFIGEKRMNDTPPAERGVGM
						VFQSYALYPHLSVAENMSFGH
1		İ				ATGLAQHKEVAEVLPSRRICWI
						ANRKRSPVVSVSVWRL
24221	54589	A	24359	364	845	RPLLDHGSSPRLNFLRFWYVVH
	1	1				AHTLMRMTEGWAGFDILRPYR
		1				WRRVIAFVVAFYFRHLIALLVV
						QMHFKLVKGNVDVVFTAQLID
						STVQLİDSTTTVTQFTHIQTQLM
					:	IQRAGTKCRKEDFRHWFVHHP
						WVL/MSPPSVTLYALFPDRSHR
		L				PRRDGSPDA
24222	54590	Α	24360	781	1275	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24223	54591	A	24361	<u> </u>	1185	MDKFDANRRKLLALGGVALG
24223	34391	^	24301	[]	1103	AAILPTPAFATLSTPRPRILTLNN
						LHTGESIKAEFFDGRGYIQEELA
						KLNHFFRDYRANKIKSIDPGLF
		1				DQLYRLQGLLGTRKPSCGYVT
						TAPAARSYRLTSKFIAVGQKAL
						SSLNIIHIAAPHLEALIIGESDTK
						NFASHDHAILIYKLEPTTGMLR
						TRAYIGQHMPLYCSAMGKIYM
						AFGHPDYVKSYWESHQHEIQPL
l.						TRNTITELPAMFDELAHIRESGR
						TMDREENELGVSCIAVPVFDIH
						GRVPYAVSISLSTSRLKQ/PRQH
						RFRLRERQPAGGTPPAGTNSVT
						GGASLHHARCQRRK*KPYRPER
						A/ESFLPARGKICWP*AGTLTVC
		1	1			IMASASRLISGCTPDSTCWSTA
						HAPICRRRGRAINRCHCSS
24224	54592	A	24362	1	801	
24225	54593	Α	24363	1523	1950	LLPA/AQKRRVDFLRATDFNAL
						MPGVVEIDGKNIYTQIIDLTTRE
						AVVNRPEVHRRYIDIQFLAWGE
1						EKIGIAIDTGNNKVSESLLEQRN
						IIFYHDSEHESFIEMIPGSYAIFFP
						QDVHRPGCIMQTASEIRKIVVK
						VALTALN
24226	54594	Α	24364	281	997	LTFARLKGITRKIGRLQSRRRKR
	1					CKAMGVDVNRVYSLDELVRG
						NDILFSATGVTGGELVNGIQQT
ļ		ļ				ANGVRTQTLLIGGADQTPMAT
		l				LTEDDVLEQLDAQDNLF\SFMK
l						TAHTILLQGIRQFLPSLFVDNDE
			-			EIVEYAVKPLLAQSGQIDDIDV
		1				ALRLIYALGKMDKWLYADITH
						FSQFWHYLNEQDETPGFADDM
						TWDFISNVNSITRNAMLYDALK
1		1				AMKFADFSVWSEARFSGMVKT

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24227	54595	Α	24365	263	1410	AMISRRRFLQATAATIATSSGFG
						YMHY/CLNSDAIALGIEQRPDLI
						LLGGDYVLFDMSLNSSAYSDV
						LSPLAECAPTFACFGNHDRPVG
						TEKNHLIGETLKSAVITGMLNQ
		1				ATVIAYANRPFELVGTGDLWS
						GQCKSASASETNLSRLVLAHNS
						CSKEVMRDEPCDLMLGSHTHR
						WQLSVPMAGEPFCHVADERYV
						LGLNAFCERHIYTIPRGGKRIPQ
		ļ				TFAADLPGKGITVNPVQSTITEE
						TFQTLLVSRALEKLGYTVNKPS
						EVDYNVGYTSLASGDATFTAV
						NWTPLHDNMYEAAGGDKKFY
						REGVFVNGAAQGYLIDKKTAD
						QYKITNIRQLEDPEIAKLVDTNG
						DGKADLTGCNPGLGFEGAINH
						QLGAYELTNTVTHN
24228	54596	В	24366	1	2872	
24229	54597	В	24367	1	1704	
24230	54598	В	24368	1	1470	
24231	54599	В	24369	1	2301	
24232	54600	В	24370	1	900	
24233	54601	В	24371	26	856	
24234	54602	В	24372	1	1743	
24235	54603	В	24373	1	1410	
24236	54604	В	24374	1	957	
24237	54605	В	24375	1	2742	
24238	54606	В	24376	755	1816	
24239	54607	В	24377	1	1968	
24240	54608	В	24378	926	6441	<u> </u>



SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24241	54609	Α	24379	1	1739	MTWTHEAVNRREPLRGCEPNW
İ	ļ					DVDVKPAGCMISRRNELPYTLG
						LRIDMLVDLGTPASLMGQSSVY
						NTTDRHSTHRMDGLEQLMQQA
						DFAADIAVVAPWDNWSPTAGR
						FGLIRLLDEQTAPYAWYTSPMH
						LMDEKCSCHRQRPVIGLNQPQ
						YQSGTIEVDNVSFAYRDDYLVL
						KNINLSVPSRNFVALVGHTGSG
ŀ						KSTLASLLMGYYPLTEGEIRLD
						GRPLSSLSHSALRQGVAMVQQ
						DPVVLADTFLANVTLGRDISEE
						RVWQALETVQLAELARSMSDG
						IYTPLGEQGNNLSVGQKQLLAL
						ARVLVETPQILILDEATASIDSG
						TEQAIQHALRRDCSPLIDHCDA
						DTILVLHRGQAVEQGTHQQLL
						AAQGRYWQMYQLQLAGEELA
		ł				ASVQPLPNSDRRGSMKLVTVII
						KPFKLEDVREALSSIGIQGLTVT
						EVKGFGRQKGHAELYRGAEYS
						VNFLPKAAYTGKIGDG/KIFVAE
	1					LQRVIRIRTGEADEAALYLWHT
						ATERDGPFRNFTAGAFRASLPH
						PARHGRRPILAAPFMPVNPSRLP
						PRRRHLASGRSLRFNCCTAGDE
						TCSARSGERRSLVF
24242	54610	Α	24380	547	684	

NO: of peptide sequence hod in USSN location of first codon for peptide sequence sequence 24243 54611 A 24381 1 2208	deletion, \=possible nucleotide insertion) MLLVSASVNAMSISGQAGKEY
sequence	MLLVSASVNAMSISGQAGKEY
	1
24243 54611 A 24381 1 2208	1
	1
	TNIGVGFGTETTGLALSGNWTH
	NDDDGDVAGVGLGLNLPLGPL
	MATVGGKGVYTNPNYGDEGY
	AAAVGGGLQWKIGNSFRLFGE
	YYYSPDSLSSGIQSYEEANAGA
	RYTICVQVFVVVMLIRNQLFTP
	VNTAIDGFATNYAIMPAYAVG
	EGVATHGLQFELMRTTGGNTQ
	HVFHHPLPLFAMWRYAAEMFP
	YQQMCQFVRYHFFNKSFTVFQ
	QQNGIEANFIAFQPGRSGGSA
	LGRNVFGFRYQPQCQQDKQYV
	QSRIHPEGVGVTQRVEHGQERC
	ANDHVQSSGTELMQEGVMSAS
	LKNQQGFSLPEVMLAMVLMV
	MIVTALSGFQRTLMNSLASRNQ
	YQQLWRHGWQQTQLRAISPPA
	NWQSGAAMLRVYHSNRLDVL
	EALMEFIVERERLDDPFEPEMIL
	VQSTGMAQWLQMTLSQKFGIA
	ANIDFPLPASFIWDMFVRVLPEI
	PKESAFNKQSMSWKLMTLLPQ
	LLEREDFTLLRHYLTDDSDKRK
	LFQLSSKAADLFDQYLVYRPD
	WLAQWETGHLVEGLGEAQAW
	QAPLWKALVEYTHQLGQPRW
	HRANLYQRFIETLESATTCPPG
	YLRASLYAAIQHALASVREHTT
	LVVIAHRLSTIVDADTILVLHRG
	QAVEQGTHQQLLAAQGRYWQ
	MYQLQLAGEELAASKGHAELY
24244 54612 A 24382 2 729	DCSPLIDHCDADTILVLHRGQA
	VEQGTHQQLLAAQGRYWQMY
	QLQLAGEELAASVQPLPNSDRR
	GSMKLVTVIIKPFKLEDVREAL
	SSIGIQGLTVTEVKGFGRQKGH
	AELYRGAEYSVNFLPKAAYTG
	KIGDG/KIFVAELQRVIRIRTGEA
	DEAALYLWHTATERDGPFRNF
	TAGAFRASLPHPARHGRRPILA
	APFMPVNPSRLPPRRRHLASGR
	SLRFNCCTAGDETCSARSGERR
24245 54613 A 24383 683 1161	RTGHCH/SAFSFSTVLTSCSLSRF
	SFLAHRLQYLLRVAEPLPNSDR
	RGSMKLVTVIIKPFKLEDVREA
	LSSIGIQGLTVTEVKGFGRQKG
	HAELYRGAEYSVNFLPKVKIDV
	AIADDQLDEVIDIVSKAAYTGKI
	GDGKIFVAELQRVIRIRTGEADE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24246	54614	A	24384	631	1149	FFGIKEQVITANNANSNRNKTD
		İ				RCCRNWLEDHADNHCHKHSEV
į	1					MPCLRRHTVGDRDKKHDTHH
		ŀ				HQWGQ*FPYIIFWLVSPLIAISRS
						KVSTHGLRQRDFILLREASQSF
			•	:		RFVLRVYDVADHIAGKFTVHD
						LQFIGYHGGQAQLVAQVFSKES
		ŀ	}			KAAGGNRHFPAQRFQLEHQLR
24247	54615	Α	24385	833	1264	
24248	54616	Α	24386	1	1632	
24249	54617	Α	24387	2078	2371	
24250	54618	Α	24388	1	159	
24251	54619	Α	24389	530	811	RPGTGTAAQVLNRRIKISRST*L
						TVVMGYWSIIPSVSLLKWDRR
						KSPKCSSALRVNISNTLLSTARR
			,			RKRYLIVIPALLVVRRCRPRGPS
			!	į		ACG
24252	54620	Α	24390	1806	2115	AAFQEIELAFQDPQCVGLLTAP
					,	MLMPLFLDQLILQCFFW/W/TN
			1			LERA*TSFSALFIDLQPPGYRTT
						TSKHKVSSSLIKGHVLLDHSFH
						DLNTQLWISTNAFRFGN
24253	54621	Α	24391	1476	1751	ALLLVSFASQRFISA/TVGNQNA
		l	ŀ			GVDLLNLTLLVGGIFVLNNAGN
						FAVFTGDTSITGRIIQFHRQQTN
		l				ATLRFSITQTLECFDRDQRHVA
						VEH
24254	54622	Α	24392	606	4519	EEVVRAISAKRNEPEWMLEFRL
	İ					NAYRAWLEMEEPHWLKAHYD
	1		ł			KLNYQDYSYYSAPSCGNCDDT
						CASEPGAVQQTGANAFLSKEV
						EAAFEQLGVPVREGKEVAVDAI
		1				FDSVSVATTYREKLAEQGIIFCS
		Ì				FGEAIHDHPELVRKYLGTVVPG
						NDNFFAALNAAVASDGTFIYVP
	1					KGVRCPMELSTYFRINAEKTGQ
						FERTILVADEDSYAAKRLTDYP
						HQLSGGERQRVMIAMALLTRP
						ELLIADEPTTALDVSVQ
24255	54623	A	24393	129	256	

SEQ ID NO:		,	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24256	54624	A	24394	1	1082	MAMSKVKSITRESWILSTFPEW
}						GSWLNEEIEQEQVAPGTFAMW
						WLGCTGIWLKSEGGTNVCVDF
ŀ						WCGTGKQSHGNPLMKQGHQM
		ŀ				QRMAGVKKLQPNLRTTPFVLD
		ĺ				PFAIRQIDAVLATHDHNDHIDV
						NVAAAVMQNCADDVPFIGPKT
ł		l				CVDSWIGWGVPKERCIVVKPG
		l				DVVQVTALEIHAPDAFDRTAPI
						TWLADQKRRRTVKGRTTRPCN
	İ					RSVTVRPRRRDAVIGRFYRHQS
		i				ITQKGLLLMVMPGCAFPLWIDS
						ELNLTFIDALFEGVSGITTTGAT
						VIDDVSSLPRAYLYYRSQLNFIG
						GLGVIVLAVAVLPLLGIGGAKL
						YQSEMP/AAI*G*QTHSPPGRYV
						TDTVDNLFFIRCRT
24257	54625	Α	24395	49	281	ASDHRYLCSQSVPLRGIPACYL
		ŀ				LQAPLTRWR/RIPPSRRICAAIDS
						AISRSPQRWLTAVCCGTVNFTN
						TCLLPDGVCA
24258	54626	Α	24396	737	1419	CIWRMYSHRLVAWRRCVKRR/
			:			RYLTRFAKTRGVAIVMVGHVT
						KDGSLAGPKVLEHCIDCSVLLD
						GDADSRFRTLRSHKNRFGAVN
						ELGVFAMTEQGLREVSNPRGR
						GKRKFISPKKTPALAKFRRNLL
						PPKVPGNGTSVRFRHSRICWMN
						PLPGAAWNCILTLKKPISPGVQ
						VFPESFTPARLEAFAGRSQACL
						VPAQNILHQKSQAWLFDFPEQE
24259	54627	A	24397	526	1693	KELRELSAWMAGA STTGIRSIAMSNPILSWRRVRA\
24239	34027	A	24397	320	1093	LCVKETRQIVRDPSSWLIAVVIP
						LLLLFIFGYGINLDSSKLRVGIFL
						EQRSEAALDF\THTMTGSPYIDA
						TISDNRQELIAKMQAGKIRGLV
						VIPVDFAEQMERANATAPIQVI
						TDGSEPNTANFVQGYVEGIWQI
						WQMQRAEDNGQTFEPLIDVQT
						RYWFNPAAISQHFIIPGAVTIIM
						TVIGAILTSLVVAREWERGTME
						ALLSTEITRT\ELLLCKLIP\YYF\
						LGMLAM\LLCM\LVSVFILGVP
						YRGSVLILFFISSLFLLSTLGMG
	ļ					LLISTITRNQFNAAQVALNAAF
						LPSIMLSGFIFQIDSMPAVIRAVT
						YIIPARYFVSTLQSLFLAGNIPV
						VLVVNVLFLIASAVMFIGLTWL
						KTKRRLD
L		L	L	L	L	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24260	54628	A	24398	655	915	IAVNVTLAFNGGKQLR*STSLL CCQEGLAFAVNLRQHFRVQTIS FFFLPVLRFWRPGIQQAKKCHP APRTAKGIPQQVIRAMRLLFK
24261	54629	Α	24399	549	827	
24262	54630	A	24400	472	626	WRFFRPRKDNIRRTLFADGHQI AAQPGLNLSCTRTIF*IKRINTFP GRWRK
24263	54631	A	24401	333	617	SDQSETDYRFADGERHLENQPH AGQTGRGGAGPGCTLYQRFAQ LCEVSEVERSGCRTLSAQHVEQ NLPAAICGGDDADGAVVHLWT TA*RTRIGTLRSGPKMNDSAISII TATDSGWKILLHMLS
24264	54632	Α	24402	74	265	
24265	54633	A	24403	1	1005	
24266	54634	A	24404	1068	1185	SVAIRQASTGNRCQCLCPACA WA*VNADVK*SARVSRL
24267	54635	Α	24405	1	2292	
	54636	A	24406	720	2496	RRAGYRKAADAMTTIVPEKLS REVGSHFHALDRTTAVHRQPVI SATTTVHEFLRCHQFPLVVCDG GRMTPRAMFLFSSAHILMPLNY HNSHINTLFPVAGTLMVEPTES ESKVELDRFIDAMLAIRAEIDQ VKAGVWPLEDNPLVNAPHIQS ELVAEWAHPYSREVAVFPAAL VTGGSRGIGRATALLLAQEGYT VAVNYQQNLHAAQEVMNLITQ AGGKAFVLQADISDENQVVAM FTAIDQHDEPLAALVNNAGILF TQCTVENLTAERINRVLSTNVT GYFLCCREAVKRMALKNGGSG GAIVNVSSVASRLGSPGEYVDY AASKGAIDTLTTGLSLEVAAQG IRVNCVRPGFIYTEMHASGGEA\ GRVDRVKSNIPMQRGGQAEEF AVQTFPVWRVAQHCAVRPFRQ RIGQLRDIFHLKGDQFTNPRKT GVTPQPPCSTYPDPSPPPVPAAG HIDRPALAPTFTPPIMCSKATVK PLSPSPPGTPDPASVLPFREQRA AVHRLRTVVTRHLLALRRDPF ANRHRQQDQRNGGFHHRQRHL HTGKTRSLHHHQFAALRQHPK
24260	54627		24407	40	196	TKQRRKQSRHREEDLHIFRHAQ
24269	54637	C A	24407	49	186	
24270 24271	54638 54639	A	24408 24409	470	811	
24271	54640	A	24410	1	1006 2793	
24212	34040	В	24410	1	2173	

sequence 09/540,217 codon for peptide of peptide sequence deletion, \=p 24273 54641 A 24411 463 1258 CCRSSWG EDRNGDG GIEHGLL HERPQKG LQGPDID GISEHVT DALVAFI MYTNPLI DAPALGI	n,/=possible nucleotide possible nucleotide insertion) CNSDSETGLQIVASYF SNLALRPEGTAGCVRA YNQEQRLWYIGPMFR GRYRQFHQLGCEVFG AELIMLTARW\WRAL LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC GIAYTVNQRLLNYRLS
sequence 24273 54641 A 24411 463 1258 CCRSSWG EDRNGD: GIEHGLL HERPQKG LQGPDID GISEHVT DALVAFI MYTNPLI DAPALGI	CNSDSETGLQIVASYF S\LALRPEGTAGCVRA YNQEQRLWYIGPMFR GRYRQFHQLGCEVFG AELIMLTARW\WRAL LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
24273 54641 A 24411 463 1258 CCRSSWG EDRNGD: GIEHGLL HERPQKG LQGPDID GISEHVT DALVAFI MYTNPLI DAPALGI	S\LALRPEGTAGCVRA YNQEQRLWYIGPMFR GRYRQFHQLGCEVFG AELIMLTARW\WRAL LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
EDRNGD: GIEHGLL HERPQKO LQGPDID GISEHVT DALVAFI MYTNPLI DAPALGI	S\LALRPEGTAGCVRA YNQEQRLWYIGPMFR GRYRQFHQLGCEVFG AELIMLTARW\WRAL LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
EDRNGD: GIEHGLL HERPQKO LQGPDID GISEHVT DALVAFI MYTNPLI DAPALGI	S\LALRPEGTAGCVRA YNQEQRLWYIGPMFR GRYRQFHQLGCEVFG AELIMLTARW\WRAL LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
GIEHGLL HERPQKO LQGPDID GISEHVT DALVAFI MYTNPLI DAPALGI	YNQEQRLWYIGPMFR GRYRQFHQLGCEVFG AELIMLTARW\WRAL LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
HERPQKO LQGPDID GISEHVT DALVAFI MYTNPLI DAPALGI	GRYRQFHQLGCEVFG AELIMLTARW\WRAL LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
LQGPDID GISEHVT DALVAFI MYTNPLI DAPALGI	AELIMLTARW\WRAL LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
GISEHVT DALVAFI MYTNPLI DAPALGI	LELNSIGSLEARANYR LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
DALVAFI MYTNPLI DAPALGI	LEQHKEKLDEDCKRR RVLDSKNPEVQALLN DYLDEESREHFAGLC
MYTNPLI DAPALGI	RVLDSKNPEVQALLN DYLDEESREHFAGLC
DAPALGI	DYLDEESREHFAGLC
	i i
	IIAYTVNQRLLNYRLS
	LSLFDGEGIYRFYHPK
	ENRQCRNDLFFRCGF
24274 54642 A 24412 221 832 TMAENQ	STVENAKEKLDRWLK
DGITTPG	GKLPSERELGELLGIK
RMTLRQ.	ALLNLEAESKIFRKDR
KGWFVT	HPRFNYSPELSASFQR
AAIEQGR	EPF\WGFTEKNRTSDI
PETLAPL	IAVTPSTELYRITGWG
ALEGHK	VFYHETYINPEVAPGF
IEQLENH	SFSAVWEKCYQKEPV
VKKLIFK	PVRMPGDISKYLGGS
AGMP	
24275 54643 A 24413 158 481	
24276 54644 A 24414 2201 2350	
24277 54645 A 24415 621 1830 AGWKRO	GAGEPDQGLQA\EME
VLENLQI	ELKDLNQRQAHAESKI
	GWFVTQPRFNYSPELS
	MEQGREPSWGFTEKN
	FLAPLIAVTPSTELYRI
	EGHKVFYHETYINPE
	QLENHSFSAVWEKCY
	KKLIFKPVRMPGDISK
	GMPAILIEKHRADQQG
1 1 1 1	YWRFEAVDLIINLCF
	ARKILQRVDTLPLYLH
	MRLERVLPADLLDIA
	VKIHVLDGERFSLGN
	SAFGDKARRLNLDIHI
	ASIDEAVAIALKTGA
	1
	RYESNLRDVLSIIAND
	YQDSRLTFNIEWGKS
	HPWAKLAAIYYRLQP
HINDAF	
24278 54646 A 24416 83 502	
24279 54647 A 24417 15 95	N 101 1 1 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1
	VPLLLQ\AYLYSAFFF
	APPLFSAAVLPEQRQR
	VIRPTVELHQALQLF
	QSLHALLRGSTGFQA
SFRLPL	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24281	54649	A	24419	1	2914	MYYSHASCVADQYACGMHRT
						PSRTGRPRIHAPRQARLKPQVK
		ŀ				VASKKRNARDKMQPPNEQQKT
1						RAHACTPSSRFRAVVRTLTCPV
ł						VANVTRIFPRYFSNTRVLRRRR
						KQARGTWWRLRHLHLYVQPH
				i		KHTSYYIYRLYRVTYVMSAQV
		İ				VDTGIYEFTTRKHDIGTIELPNS
						EIRYHGNPAMSRPRKTPTPPNSP
			}			ASERATGPTTLLPPIRQRRYAA
						RHTSTIAYTNSTSDIFEAPYNKA
						APIMRYTSVYCSLFSP
24282	54650	Α	24420	2	420	CPYPENRWRGKILARPLLFRAA
ł					1	*ASRFAGAGMPGRLIPTVRPRN
ł						SNSGRLVSASRCAKESLTVTTC
						TPLPGSAVREASN
24283	54651	A	24421	212	497	
24284	54652	Α	24422	1	1149	
24285	54653	Α	24423	535	589	
24286	54654	Α	24424	1	1458	MFDPETLRTFIAVAETGSFSKA
ŀ		l				AERLCKTTATISYRIKLLEENTG
						VALFFRTTRSVTLTAVGEHLLS
						QARDWLSWLESMPERAATVW
1						MFVPTPLSVASGLQRIGRAGHQ
						VGGVSKGLFFPRTRRDLVDSAV
						IVECMFAGRLENLTPPHNPLDV
	1					LAQQTVAAAAMDALQVDEWY
						SRVRRAAPWKDLPRRVFDATL
ŀ						DMLSGRYPSGDFSAFRPKLVW
						NRETGILTARPAAFAADSDIAD
						GQTQRFDFSILQSMAHDLSQTA
						WR/GIQYDAEKSLWHNVE\NRQ
İ						LDAQFFHMGMGFRRRVRMFSV
						DPATHLAREIHFRPELFKYNDA
						GVDTKQLEGQSDLGFAGFRVF
1					ŀ	KAPELARRDVVSFLGASYFRAV
1						DDTYQYGLSARGLAIDTYTDSK
						EEFPDFTAFWFDTVKPGATTFT
						VYALLDSASITGAYKFTIHCEKS
1						QVIMDVENHLYARKDIKQLGIA
1		l	}			PMTSMFMPRKNASRLSKRCYDI
		<u> </u>				RERDGREERK
24287	54655	Α	24425	157	650	PIRASTCLFKPLITQCFRQFGSE
						GVNDHLLGILLIDPARAQIEYLF
					1	IINTPHRRAVAAFHIICIDFQLRF
						RIHFRQFAHQQIVVGHLTIRFDG
						VLRDVDQTVKYRAAFVADDSF
						VQLSAVTKTFIVFQPGTGITHLV
						FHRHRQAQ*PFKVPFSRRLRNGI
1000		 	2442	2.0	2024	НТНС
24288	54656	Α	24426	269	2024	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1		Ì	sequence		
24289	54657	Α	24427	1	3548	MPRSNYFSERCKGEIFLKFENM
2.205	3.037		21127			QRTGSFKIRGAFNKLSSLTDAE
						KRKGVVACSAGNHAQGVSLSC
						AMLGIDGKVVMPKGAPKSKVA
						ATCDYSAEVVLHGDNFNDTIA
						KVSEIVEMEARIFIPPYDDPKVI
					i	AGQGTIGLEIMEDLYDVDNVIV
						PIGGGGLIAGIAVAIKSINPTIRVI
				i		GVQSENVHGMAASFHSGEITTH
						RTTGTLADGCDVSRPGNLTYEI
				! !		VRELVDDIVLVSEDEIRNSMIAL
						IQRNKVVTEGA
24290	54658	В	24428	1	2691	
24291	54659	Α	24429	2174	3315	NLPKPDWARSAPHPASANYKS
						NTRRLPFAREQ/QDQHTLLAGT
						LFIAKILIPMQIAQSAAYRRRCP
						DAQLHFLPLQLAPRGHRYKRV
İ						CQVRYPPIIALIPVVFRLMNDAN
l		ŀ				GLSQPADAGLAHHEMRNFLDF
						ASSNVAASRRRCVAALPCATG
						KGRQANEQAENKCSKDPQTHR
				ļ		TDNKETKTPPARPTTTTQPHTK
						EEEPLSTPTQTKHPPTHVTGTGP
						HPTPASGQTEKHERKRPTQQKS
						NARQNSKDKRTPNKSKTTKPN
						RGYNQHPAGEPTMSTHEKVKP
						HQQNRKIQAPPKRQKTDHRHT
						HNTHPHPPHRAASTTTQAAPPP
						RRATRSGTSATAHNNTEPHGTR
						HTQPGDHKRAEEHDRTADEPT
						PPEHTLPSHTQLGP
24292	54660	Α	24430	1	2925	
24293	54661	Α	24431	341	689	RPLAFRQRCTFYWR\PGSANCR
						LV*GASGTDPRLYSPCAAAADD
						CGRRQNAGRRRRATSGD*SPHR
						RWENALLFDSRHRHCPRRAKN
						AGGEYRQRGIAGSDLQQRFTA
						AEKDHSRS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 ' ' '
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24294	54662	Α	24432	1	1931	MGTLLGTLTVLTLVAEELPAET
						DVAADPAASPTCTAVNQTIGRA
						IVAADFFIIFQLWQNTVRQLVT
	1					HFKPHGQSEMSDTPELINRARY
						LTYLYCRQPAKAIRELSEQELA
						WKKQATHEYQVLVTRCKQTL
		İ				HQVEPLREIEPGRKRLVFDENL
						RPKQNPIPHLLDTCWPLTVDEH
						ANRSSIGNNNISFVVNRAVVTN
						VLNTMRTACNSQATIQIYNADI
						ARDFGTRGIFSINSGFSTVIRGG
						VVTVDLSAGAAIGDSALAFVID
ļ						DGLVAVNLDTDGLRIGDGHGA
						LVIHRCAVAGDINTVTAAADM
]						DLEDPRIISARFGVPRTQVRTW
						VALYEKHGEKGLIPKPKGVSAD
						PECVLGRESCDRAAHVPYQAA
						AHFMLAGSGSVARWLKVYEER
						GEAGLRALKIGTKRNIAISVDPE
						KAASALELSKDRRIEDLERQVR
		ļ				FLETRLMYLKKLKALAHPTKK
ļ						AAEIPRSTFYYHLKALSKPDKY
						ADVKKRISEIYHENRARYGYRR
1						VTLSLHREGKQINHKAVQRLM
1						GPSHLKQRLRSSDTALTEERDF
						KATRPNEKWVTDVTEFAVNGR
						KLYLSPVIDLFNNEVISYSLSER
						PVMNMRRSRPGVGYFYALTPV
						YLANRLQALLGVEQ/HHPDGEL
						FLTGYVADGGYHS
24295	54663	Α	24433	3	158	
24296	54664	Α	24434	1	3432	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24297	54665	A	24435	1298	2649	TFITRPTSNGTSIFKLMNRRRLE
						GGNVATLR/RLTIGMIFQDHHL
						LMDRTVYDNVAIPLIIAGASGD
		1		-		DIRRRVSAALDKVGLLDKAKN
						FPIQLSGGEQQRVGIARAVVNK
	1					PAVLLADEPTGNLDDALSEGIL
	1		Ĺ	1		RLFEEFNRVGVTVLMATHDINL
						ISRRSYRMLTLSDGHLHGGFGG
						RLDRFRKSVGGSGDGGRNAPK
				•		RAKSSPKPVNRKTNVFNEQVR
1						YAFHGALQDLKSKPFATFLTV
]	MVIAISLTLPSVCYMVYKNVNQ
]		AATQYYPSPQITVYLQKTLDDD
}	1					AAAGDPDRPSLQLRWRVIPGFS
					•	IAVAIYFLNESLGRPLMITGYSA
						ECSLLAWPVRSGLRKAPMKTFP
						LQSLTIIEAQQKQFALVDSICRH
		i				FPGSEFLTGGDLGLTPATELKPR
						RLPSGVEQVLADAFHAQAAAL
						VQGAGTGAIRAGLAALLKPGQ
						RLLVHDAPVYPTTR
24298	54666	A	24436	546	707	ICEVIIDA VIIIA
24299	54667	A	24437	428	1464	RNSRSLAEISRLRGQTRSGLPM
24277	34007	`	24437	120	1101	LLNLQSMGAS/WYLSPVIDLFN
		1				NEVISYSLSERPVMNMDHHLL
1						MDRTVYDNVAIPLIIAGASGDD
1		1				IRRRVSAALDKVGLLDKAKNFP
						IQLSGACWHCRAVVNKPAVLL
1						ADEPTGNLDDALSEGILRLFEEF
	1					NRVGVTVLMATHDINLISRRSY
		l				RMLTLSDGHLHGGFGGRLDRF
1						RKSVGGSGDGGRNAPKRAKSS
		1				PKPVNRKTNVFNEQVRYAFHG
		1				ALQDLKSKPFATFLTVMVIAISL
		1				TLPTSCYMVYNIAKSERISDIST
		1				OLNAFPGCEVAVSDAPSGOLIV
		1				VVEAEDSETLIQTIESVRNVEGV
					:	LAVSLVYHQQEEQGEETP
24300	54668	A	24438	1	429	EAVSEV ITIQQEEQGEETI
24301	54669	A	24439	307	424	
24302	54670	A	24440	300	468	
24303	54671	A	24441	946	1558	KGCVSIPFFSFSWAVMSNLTYL
[,,,,,,	37.7	Γ.	I	1		QGYPEQLLSQVRTLINEQRLGD
		1		-		VLAKRYPGTHDYATDKALWQ
		1		1		YTQDLKNQFLRNAPPINKVMY
						DNKIHVLKNALGLHTAVSRVQ
		1			}	GGKLKAKVE\IRVATVFRNAPE
						PFLRMIVVHELAHLKEKEHNK
		1				AFYQLCCHMEPQYHQLEFDTR
					1	LCNRIRLNGHDHFRRRYALFRV
		1			1	NTGRTASPVIQP
L	<u> </u>	<u> </u>		<u> </u>	1	1410/11/101 /101

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24304	54672	Α	24442	1	774	
24305	54673	Α	24443]	452	MELKKLMEHISIIPDYRQAWKV EHKLSDILLLTICAVISGAEGWE DIEDFGETHPDFLKQYGDFENG IPVHDTIARVVSCICPAKFHESFI NWMLDYHSSDDKD/DHRN*W KNTPAFL*QESP*GSDSCH*CVL NNAQSGHRTDQDG*EIQ
24306	54674	Α	24444	486	572	
24307	54675	A	24445		683	MHSLVIGQIKTDKKSNEITAIPE LLNMLDIKGKIIKTDAMGCQKD IAEKIQKQGGAVMNLQRFDDST LIRIFALHELHRLKEHGLTRGAL LDYHSRYKLVFL\AHPQPESRK LGPF\VADIHQWQNLDDYSPQY RQRVVVLLSHPANARDHTNVL MHVQGYFRPHIDSTERQQLRAL IDSYRRGEQPLLAPLMRIKHYM ALYPDAWLSGQRYFELWPRVI NLRHSGVL
24308	54676	A	24446	896	1066	WRLSPSSR*LMPKVPRIQAYFT T*YCS*VIQRSPC*FHCYSRSIPW AWGAGKQSLT
24309	54677	Α	24447	1	3660	
24310	54678	Α	24448	1	1530	
24311	54679	Α	24449	1	1890	
24312	54680	Α	24450	746	1872	KHWRGSLISLSTPARFNGGPFW GANQGGDPKGRGGSQRALFRR KNNRGGETVSGAEDACIVNNN AAAVLLMLAATASGKEVVVSR GELVEIGGAFRIPDVMRQAGCT LHEVGTTNRTHANDYRQAVNE NTALLMKVHTSNYSIQGFTKAI DEAELVALGKELDVPVVTDLG SGSLVDLSQYGLPKEP\MPHELI AAGVSLVSFSGDKLLGGPQAGI IVGKKEMIARLQSHPLKRALRA DKMTLAALEATLRLYLHPEALS EKLPTLRLLTRSAEVIQIQAQRL QDPLAAHYGAEFAVQVMPCLS QIGSGSLPVDRLPSAALTYTPH DGRGSQLESLADRWRELPVPVI GRIYDERLWLDLRCLEDEQRFL
24313	54681	Α	24451	1358	1419	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	•	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24314	54682	A	24452		1568	MKGRNIDAQEIQKRYGMVGDP VIIGVVLGLIFGLAAGEGFKGC ASLMITVAAIMVLFPRMIRLIVE GLLPISDGARKFFQKYFKGREV YIGLDTAVTLGHPTTIAVGLLLI PIMLILASILPGNKVLPLADLPV APFFICMATVIHRGDLVRTLISG VIVMITVLLIATQFAPYFTEMAL KGGFSFAGESAQISALSVGNMF GWSISELMSLGIIGRVVAVGIVA SVVLFLRVLSMSWLKEVIGTEK AVIAMCHLRALPGDPSFDAQLG MNWVIDKAWDDLMALQNGG VDAVMFSNEFSLPYLTKVRPET TAAMARIIGQLMSDIRIPFGVNV LWDPVASFDLAMATGAKFIREI FTGAYASDFGVWDTNVGETIR HQHRIGAGEVKTLFNIVPEAAV YLGNRDICSIAKSTVFNNHPDA LCVSGLTAGTRTDSALLKRVKE TVPDT\VVLANTGVCLENVEEQ LSIADGCVTATTFKKDGVFANF VDQARSNNPSAIPAAFVKSIDG DRYQINPNRHGVNDG
24315	54683	В	24453	1	3352	2111 Q
24316	54684	В	24454	60	2284	
24317	54685	В	24455	1	2475	
24318	54686	В	24456	1	2890	
24319	54687	Α	24457	1065	1139	
24320	54688	Α	24458	282	615	
24321	54689	Α	24459	2657	2919	
24322	54690	Α	24460	1	835	
24323	54691	Α	24461	167	338	
24324	54692	A	24462	834	1515	LRWCSMPCCRWSKDRAAPSPA YSRLSSHVIIAGYVGGGVARG\ HSWVFVALTLVLTAVLFSLAGL LNGVFAKTFDDISLVPTFVLTPL TYLGGVFYSLTLLPPFWQGLSH LNPIVYMISGFRYGFPWLNHNR PGVAVDINRLLLPRKQEKLGIK LDKDQLIAGVQDAFADKSKLS DQEIEQTLQAFEARVKSSAQAK MEKDAADNEAKGKEYREKFR QIPSVSSCQKQ
24325	54693	Α	24463	3	557	
24326	54694	A	24464	255	693	
24327	54695	A	24465	3	242	QKAPVLANQRQPRRPTGALLLE /SLPLVAESSDG*EPMTNATEPH SRPLP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleofide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24328	54696	A	24466		2636	MKNIIRTPETHPLTWRLRDDKQ PVWLDEYRSKNGYEGARKALT GLSPDEIVNQVKDAGLKGRGG AGFSTGRKWILMPKDESMNIRY LLCNADEMEPGTYKDRLLMEQ LPHLLVEGMLISAFALKAYRGY IFLRGEYIEAAVNLRRAIAETTE AGLLGKNIMGTGFDFELFVHTG AGRYICGEETPPLLKPQTGVWQ RHSCWKLQELAMNDRDFMRY SRQILLDDIALDGQQKLLDSQV LIIGLGGLGTPAALYLA
24329	54697	Α	24467	506	673	
24330	54698	С	24468	1	3411	
24331	54699	A	24469	2455	2694	
24332	54701	A	24471	02	1081	MGGLRDPWLHKLRSLWGLCR YGHKEQYDHIRIELNLAPGDTT LVPTGLAIHIADPSLAAMMLPR SGLGHKHGIVLGNLVGLIDSDY QGQLMISVWNRGQDSFTIQPGE RIAQMIFVPVVQAEFNLVEDFD ATDRGEGGFGHSGRVFCNMAE KQTAKRNRREEILQSLALMLES SDGSQRITTAKLAASVGVSEAA LYRHFPSKTRMFDSLIEFIEDSLI TRINLILKDEKDTTARLRLIVLL LLGFGERNPGLTRILTGHALMF EQDRLQGRINQLFERIEAQLRQ VLREKRMREGEGYTTDETLLA SQILAFCEGMLSRFVRSEFKYRP TDDFDARWPLIAA\SCSNMTPD
24333	54701	A	24471	93	475	RCKCADAATRGKSRRDLPDVQ NSARQYG/WI/GCATHARSG*HI VEAVQSG/TPPQPLRPRGTSRKV STDRYWRAASQLPAQYQDLTSI YDVYTVPDRVAKRAFKSPPDD GPVMWRMGHQPHLIPRNVLAQ
24334	54702	A	24472	254	1150	RFLTCRYGPGMAEDMQSLVGG TVTRSG*LSVVVNNNGE/LLAK ELSVQVGIRPEYLDRLLHAFSG GQRQRIAIARALSSQPDVIVLDE PTSALDISVQAQILNLLVTLQEN HGLTYVLISHNVSVIRHMSDRV AVMYLGQIVELGDAQQEIAQA LRRTPAPMVYIGNLGRELSLPA ANLKLESKLAIMEQYVGKKVID AVIVGPKVDVSAVKERIVIQEV LEASDIPYRHDRQLLHNALEKA LQALGGLEVIQLYRRDSNLVPG QLLKTSEMGKTRLYAADSRLIA AKGSPIQPTLDSLK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24335	54703	A	24473	1874	2927	KEDQVKGGMVCPKKLLCQGTR
						*RGGVVGGISQASVTSVIGYSA
						SCQPTPAIMQLHL/ANAASRRL
	1				1	QPDRNRPNPGWHRPPIHAFTTP
						GLVSVSFLGYPCSTPTCLPFSES
					† 	SVGESVNLWPQSNENLRRTLV
						GEGNLLLTFIGNGKHEIIASIFFA
						FSEGISAQSHWQPRYISPASVCT
						LVTEIDIKANKITFSIFDENAGA
						WLPPFCKAVVPGWVYKPNLGH
						EIACVDHWLMPNRLIENLCDIM
ł						LPIWVLTKFNDCCAGCVPCCVII
1						VDDALHGAPKRMCCCITRGMR
						LLKLSVHAATRSTLQQASHGH
					,	CAWSAPGWLDTSGMDGDRLW
						WTLHEVLRDLCTSHGNIVESAR
24336	54704	В	24474	1	672	
24337	54705	В	24475	1	2496	
24338	54706	Α	24476	3260	3363	RPDSGGP*RQRRYLRCADHRRR
						AVRNQQQLTACA
24339	54707	Α	24477	112	224	
24340	54708	Α	24478	1614	2326	CTYSRKCRPDKRSASGNFALSS
]						LKSSPLTKNRRYWKVSPTHLLL
						SKRPKISLNAYFGACSLVRPVL
						AAASDWTLQGAWSPTRAKALP
						ICESWRIPYADSLSSLAASCDAV
						FVHSSTASHFDVVSTLLNAGVH
						ICVDKPLAENLRDAERLVEG\M
						REWREERGQGVVHKPIPGWQS
						TLEQRGFVGCARHFIECVQNQT
						VPQTAGEQAVLAQRIVDKICSS
						FNVVRVAPLGDPIQSKPVV
24341	54709	Α	24479	19	885	
24342	54710	В	24480	1	1704	
24343	54711	Α	24481	974	1156	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24344	54712	Α	24482	162	1602	HRRKRARGTPKATTTRKPARTI
						ETLSPLLAILAFAPKLAICTRSPL
						L/DDMTPFTQCTNAYSHSGIRNP
						TAISSAAYPRRITGY/KLGPTGE
						ERTVYGIYGISESNLAECEKGV
						KSAVALTPALQPIDGVAVSYID
						AAVALGNTINEMDKYYTQENY
						KDDAFAKGKTLHQTFLKNLEA
						FEPVAESYHAAIQEINDKRQLA
						ELKNIEEREGKTFHYYSLAVMI
						SAKQINNLISQDKFDAEAAMKK
						VSELETLVAQAKEADKGGMNF
						SFINSAGQYQLEAKKYVRRIRD
	1					` `
						KVPYSDWDKEQLQDANSSWM
						VEDSFPRALREYNEMLMTRIGG
	1					GAGDQDMVNDFSYSPQLVGRG
						WFNDQVATGCTRDSTFNQQQV
	}					ALSVNAYYVVNRFGKIDWKPM
	1					YAEADKLLAKPNMRFKSDKLV
ļ						GDLSIGDQQMVEIAKVLSYESK
	1					VIIMDEPTDALTDTEAESLFRVI
24245	54712	 	24402	ļ	10.62	REMKSQAPRYCLYLPPHENNL
24345	54713	В	24483	1	1263	
24346	54714	В	24484	148	1071	
24347 24348	54715 54716	В	24485	168	1088	
24348	54717	В	24487	1	1236	
24349	54718	В	24488	1	849	
24351	54719	В	24489	1	1548	
24351	54720	В	24490	1	1372	
24352	54721	В	24491	1	2469	
24353	54722	В	24492	1	1917	
24355	54723	В	24492	1	1159	
24356	54724	В	24494	1	2294	
24357	54725	В	24495	217	2476	
24357	54726	A	24496	477	916	SLPECEDRVQYRRSLFADPSYR
24336	34720	^	24470	17''	910	YGLARHGDPRRAGRLRDV*PL
						HE*YRHWHDQS/RLLIASVLPQ
						VWHGACNAMTSSCIARQVS*N
1						`
1						TALIIRMRKGDLSAVMSSQTGL
1						INIVRRGLLLSCFRLTAMKISTV
04050	54707	ļ	24407	1126	1217	SPFRPQMPSIVRSVWC
24359	54727	A	24497	1136	1317	
24360	54728	Α	24498	3	317	
24361	54729	Α	24499	1	1245	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24362	54730	ĪA	24500	85	1293	ITVATTQHQAGLTLRRAGIRQP
						DMAVQRDHLPFRVAGVIRHIM
	•	l				LMLHSQRRTGLLITAKNRLLFF
į				1		QRRSVPKFHSKLNQAYCLIAAT
						DCPRRLLLICRYPHHFLHQLIIL
						AIGIGSAVFEPHQIIGVSRCCPR
						HLGVERVLFLFLNDFIEQVIIRC
ŀ						HVVCQSAPAAANIQYAHPWF*
			į			LQFFTNHFELRLLGRFQIIGIFPV
		l				SAGILHEGIKHAAEQVITQIVML
		į .				FADYPGTFFTL/RD*RDERRQY
						AACL*NDGGVGSRDLRVRLC*
		İ	İ			KTYPVGHIPTSHPYN/CSPNPKE
						PSCRTRLKKSGFFTRISAGAEPL
						MAIPARLSNSLACGARDSAAKE
		İ				VLLWFRFVLLLKQTLLNWTTK
			ļ			NVTSLWRTLGSPVGVKEVRAW
						TIPV/VSNRAAGSGQNPY*F*KR
24363	54731	Α	24501	2794	3531	SPRDSPTFWTKGGPGGSRGKAL
						RLVVGTDFHQAADNIPDHMMY
						KSIRFNIYHHIRTVTRDINMHNV
						APRGSGLALHRTERGEIILAQQ
						ALRSPMHPFSVQRFVKMGYPFT
						QYRRAQPMIVDDVAITASRGA
						KSGMKIIRYSFNPTHSDVTR*IRI
		1				YAQGPRRVFAQRGCFEMHNLP
						GTVYPCVSTPGAEDGDGFVRH
						LGERLFQLLLHTADFVLPLPAVI
24264	54722	-	24502	1	1767	LTGGVLNAQRNFINRRLLPYVE
24364 24365	54732 54733	A	24502 24503	1	1275	SNPTAIGGRPRRPYQAEDRSLD
24303	34733	A .	24303		1273	FVDTALNRKILKDSIFHALQTIM
						VGIKDFLRLTQVFFDLATRIPRH
						LYHPVDITTHYGRFCRHRRHHF
						QLLQLCFRFLFCLFRHLRRVDF
						ALQGFVFVRRVVHFTEFFLNSF
			ľ			HLLVQIVLTLGFLHLLFNAVAN
						AFLNLQQIDFRFHHCHQIFQTFV
						NVGHLQNRHRKIGIVIFRSDVL
						EHPHRNNTIKLVVQVTIILQQN
						GDIQAFTAFLSHFLLFGRNGDA
						HHAYVIIRCHVVCQSAPAAANI
						QYAHPWF*LQFFTNHFELRLLG
						RFQIIGIFPVSAGILHEGIKHAAE
						QVITQIVMLFADYPGTFFTLQIE
						ETSAGNTQRVFEMMGELVLET
						RLRSPAGFTQWIPHQGCRWSCL
						PVPRGTLALLSPWVVDGTGRR
						GAGGGTPQGGSSCTGAHGGGG
24366	54734	Α	24504	1	2052	
	1	1	1	<u> 1 ' </u>		L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24367	54735	A	24505	11	1060	MNARQQVAALRVPGVEKATIV
	.,,,,,					YRRSLQEMPAWREEYEEALHD
				1		GFRFLNNPERFDADGTLTLRVM
				}		SLGEPDEKGRRRPVETNETVTL
						LVDSLITAIGEQQDTEALNAMG
						VPLDKNGWPDVDHNGETRLTD
ŀ						VFMIGDVQRGPSSIVAAVGTAR
İ						RATDAILSRENIRSHQNDKYWN
						NVNPAEIYQRKGDISITLVNSDD
İ						RDAFVAQEAARCLECNYVCSK
						CV\DVCPNRANVSIAVPGFQNR
						FQTLH\LEA\YCNECGNCAQFCP
			İ			WNGKPYKDKITVFSLAQDFDN
						SSNPGFLVEDCRWVPHRGLYH
i.						RLYREPTWRSEWPRRHICPEGS
	a de la companya de la companya de la companya de la companya de la companya de la companya de la companya de					TFPERHCDHLSANNSRNKLLPL
;						OKPGFIN
24368	54736	A	24506	1316	2678	QKIGIIIV
24369	54737	A	24507	2172	5242	IYFSERLPALCTPDAHYREPCQ
2-1307	34737	' '	24507	2172	3212	YNCTRLDYDSALNIRELKKVAL
						EKGWDEYKQRWHKPAGSGSR
						HPVAVIGAGPAGLAAGYFLAR
						AGHPVTLFEREANAGGVVKNI
1						VPQFRIPAELIQHDIDFVAAHGV
Ì						KFEYGCSPDLTIEQLKNQGFHY
1				1		VLIATGTDKNSGVKLAGDNQN
1						VWKSLPFLREYNKGTALKLGK
1						HVVVVGAGNTAMDCARAALR
1						VPGVEKATIVYRRSLQEMPAW
		1		1		REEYEEALHDGVEFRFLNNPE
24370	54738	A	24508	192	484	RGLVHRGRYRWRQKWSAGSP
24370	34730	'	24300	1172	101	ARHPV/SQSTTDTPATHALSTKII
						SPIPNAPISALARTTATRL*RSM
						NQPPVKRPTVIPRVKPA*ISEPE
						VGPIILGP
24371	54739	A	24509	1	1713	V GT ALLOY
24372	54740	A	24510	1	651	
24373	54741	A	24511	3	783	PVLSKTPLTAKAIDAAQPQDKP
12 15 15						YKLTDSLTPGLFLLVHPNGSKY
						WRFRYWLNKREFLQAIGVYPLI
						TLKEARRATESRSLIANGINPV
						EQARKEKAIDALNMAAGFKKV
						AEDWFATRVGGWSESYAKQV
						RSALEKDVYPVLGKRSIVDITA
						RDVLALLQKKERTAPEQARNV\
						AGASVRSSSLPLSPNWGGWLA
-	İ					RAAFPHHITIAARMTKHPPCW
						WCTILACRQASLAVRGSTHYSL
						ELLIRRHILSLLRSPICASPLTV
	<u></u>		L	<u> </u>	<u> </u>	ELLIKKHILSLLKSPICASPLI V

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24374	54742	IA	24512	746	2387	WTALPARLKQIRSGRKLPLTVM
			,			PALRETDTFDTFMESSWYYAR
						YTCPQYKEGMLDSEAANYWLP
						VDIYIGGIEHAIMHLLYFRFFHK
						LMRDASMTNPSSVPRAGGPAL
						LFENPKGYSMPVLCNLFGTPKR
		1				VAMGMGQEDVSALREVGKLL
		1		-		AFLKEPEPPKGFRDLFDKLPQF
						KQVLNMPTKRLRGAPCQQKIV
						SGDDVDLNRIPIMTCWPEDAAP
						LITWGLTVTRGPHKERQNLGIY
						RQQLIGKNKLIMRWLSHRGG\A
						LDYQEWCAAHPGERFP\VSVAL
						GAVPATILGAVTPVP\DTLSEYA
			ļ			FAGLLRGTKTEVVKCISNDLEV
						PASAEIVLEGYIEQGETAPEGPY
						GDHTGYYNEVDSFPVFTVTHIT
						QREDAIYHSTYTGRPPDEPAVL
						GVALNEVFVPILQKQFPEIVDFY
						LPPEGCSYRLAVVTIKKQYAGH
						AKRVMMGVWSFLRQFMYTKF
]					VIVCDDDVNARDWNDVIWAIT
						TRMDPARDTVLVENTPIDYLDF
						ASPVSGLGSKMGLDATNKWPG
		1				ETQREWGRPIKKDPDVVAHIDA
24375	54743	A	24513	576	878	NLGSVWHYFRRSPEPATHHDA
						ANLERSPAA*RLSGARYRILAV
						*ADQSQTGSGDGSPDLQTGRV
	}	1				GDEARHRKRGLHVPQPRSEPPV
1						GARGFPYRFATRWR
24376	54744	Α	24514	2	878	
24377	54745	Α	24515	1	735	
24378	54746	Α	24516	1	1992	
24379	54747	Α	24517	721	1239	

3220

24380 54748 A 24518 I 1954 RQLNGVQLAFRESFHTVTRVT QHAAGAVAIHQHIRNQLITGAI GIFAVAVCLQQRLDILLAGH GIFAVAVCLQQRLDILLAGH GIFAVAVCLQQRLDILLAGH AEHVQFVIAQFVTCQQHGNGN RNTIIFILFNELRKIVETVRVC QTQTGTGEVAFHPQLFRGRVVY PDLNARRGNGVINCSTLEDPO GLILFVYTIGIQVGPGFFASLRX SGLRLNLFAVLIVIIGGLVTAIL HKLFDIPLPVVLGIFSGAVTNT ALAGAGQUICRDLGTTMEND MGMSYAMAYPFGICGILFTIM MLRVIFRVNVETEAQQHESSR NGGALIKTINRVENPNLHDLA KDVPILNGOKIICSRLKREETLI VPSPDTIIQLGDLLHLDSRGF DTQDIDLQNSSVEKQFGDAIR TILDVLARYNVRGVQLNVDDI GALDCILRARLEALLARAGE AGE GALDCILRARLEALLARAGE AGE GALDCILRARLEALLARAGE AGE GSTGLLAALESPLGITTAVELAH HPLYRDIETIVRVNALDSEWG NDLEAVVRGGADVRITKTD AQDVLDIEKEILRIEKACGREF GSTGLLAALESPLGITRAVELAH ASERLIGIALGAEDYVNRLTKTD AQDVLDIEKEILRIEKACGREF GSTGLLAALESPLGITRAVELAH ASERLIGIALGAEDYVNRLTKTD AQDVLDIEKEILRIEKACGREF GSTGLLAALESPLGITRAVELAH ASERLIGIALGAEDYVNRLTKTD THE RSPEGTELLSARCPHWQAARSA GIQAFDTVYSDANNEAGFL/QI LYFTAISYGKULVGPFBAGANAMVNSST TFGIVSGISDGHVPASYLLALA VLFTAISYGKULVGPFBAGANAT TFGIVSGISDGHVPASYLLALA VLFTAISYGKULVGPFBAGANAT TFGIVSGISDGHVPASYLLALA VLFTAISYGKULVGPFBAGANAT TYAQKSINPHVGFMYGWSSLL DCLLAVILAAIYVGFLAGFH YIGMLFVFGANAAMVNSSL TYAQKSINPHVGFMYGWSSLL DCLLAVILAAIYVGFLAGFH YIGMLFVFGANAAMVSSL TYAQKSINPHVGFMYGWSSLL DCLLAVILAAIYVGFLAGFH YIGMLFVFGANAAMVSSL TYAQKSINPHVGFMYGWSSLL DCLLAVILAAIYVGFLAGFH YIGMLFVFGANAAMVSSL TYAQKSINPHVGFMYGWSSL DCLLAVILAAIYVGFLAGFH YIGMLFVFGANAAMVSSL TYAQKSINPHVGFMYGWSSL DCLLAVILAAIYVGFLAGFH YIGMLFVFGANAAMVSSL TYAQKSINPHVGFMYGWSSL DCLLAVILAAIYVGFLAGFH YIGMLFVFGANAAMVSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSINPHVGFMYGWSSL TYAQKSI	SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
QHAAGAVAIHQHRNQLITGAI GIFAVAVCCLQRCIDILLADHI AEHVQFVIAQFVTIQQHODILLADHI AEHVQFVIAQFVTIQQHODILLADHI AEHVQFVIAQFVTIQQHODILLADHI RNRTIIFLLFNELRKIVETVRVQ QTQTGEVAFHPQLFRGRVVYC PDLNARRGNGVINCSTLEDPQI GLILFYYTIGIQVGPGFFASLRI SGLRINLFAVLIVIIGGLYTAIL HKLFDIPLPVVLGIFSGAVINTI ALGAQQQILRDLGTPMEMVDI MGMSYAMAYPEGICGILTATI MLRVIFRVNVETEAQQHESSRI MGASALIKTINIRVENPINLHDLA KDYPILINGRIGISKILKERI VPSPDTIQLIGDLILHLDSRGSPI DTQDIDLQINSSVEKÇFGDAIR TIILDVLARYNVRGVQLINVDDI GALDCILRARLEALLARASGLI MISASLQQRKTRTRRSMLFVPF ANAAMVNNSTIYADALMFDI EDSVALREKDTARRMYYIIAL HPLYNDIETIVRVNALDSEN HPLYNDIETIVRVNALDSEN ANAAMVNNSTIYADALMFDI EDSVALREKDTARRMYYIIAL HPLYNDIETIVRVNALDSEN ASERLIGIALGAEDYVRNLRTE RSPEGTELLSARCHWQAARSA GIQAFDTVYSDANNEAGFLQQI ASERLIGIALGAEDYVRNLRTE RSPEGTELLSARCHWQAARSA GIQAFDTVYSDANNEAGFLQQI ASERLIGIALGAED VRIPKTBIAGAPI A 24520 2520 3525 A 24521 1 1263 MAINSPLNIAAQPGKTRLRKSI KLWQVMMGLAYLPMTVFI TFGIVSGISDGHVPASYLLALA VLTTAISYGKLVRQFPEAGSAN TYAQKSINPHVOFMYGWSSLL DCLLAVVILAIVYGFSLAGH VVRLYENDTAQDVLDIEKEILR EKACGREPSGTIGLALASSTOTI RMVYHALGHPLYRDIETIT TRAVEIAHASERLIGIALGAED VRNLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALAGSED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALAGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALAGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALAGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALAGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLALGAED VVRLPKTDTAQDVLDIEKEILR EKACGREPSGTIGLAGAED VVRLPKTDTAQDVLDIEKEILR EKACGR				031010,211	1		,
SGLRINFAVLIVIIGGLYTAIL HKLFDIPLPVVLGIFSGAVITI ALGAGQQILRDLGTPMEMVD MGMSYAMAYPFGICGILFTMY MLRVIFRVNVETEAQQHESDLGTFMEMVD MGMSYAMAYPFGICGILFTMY MLRVIFRVNVETEAQQHESDL MIGSALQGILRDLGINSVEKGFGDAIR TILDVLARYNVRGVGQLNVDDI GALDCILRARLEALLARASGLI MISASLQQRKTRTRRSMLFVPP ANAAMYNNSFITYADALMFDI EDSVALREKDTARRMVYHALI HPLYRDIETIVRVNALDSEWG NDLEAVVRGGADVYRTIKTD AQDVLDIEKIELRIEKACGREF GSTGLLAAIESPLGITRAVELAH ASERLIGIALGAEDYVRNLRTE RSPEGTELLSARCPIWQAARSA GIQAFDTVYSDANNEAGFL/QI 24382 54750 A 24520 2520 3525 24383 54751 A 24521 I 1263 MAINSPLNIAAQPGKTRLRKSI KLWQVVMMGLAYLTPMTVFI TFGIVSGISDGHVPASYLLALA VLFTAISYGKLVRQFPEAGSAY TYAQKSINPHVGFMVGWSSLL DCLLAVVIIAAIYVGFSLAGFH YIGMLFYPGANAMYSNSFIY ADALMFDLEDSVALREKDTAI RMVYHALQHPLYRDIETIVRV ALDSEWGVNDLEAVVRGGAD VVRLPKTDTAQDVLDIEKEILB EKACGREPGSTGLLAAIESPLG TRAVEIAAMSPRIJAGAGD VVRLPKTDTAQDVLDIEKEILB EKACGREPGSTGLLAAIESPLG TRAVEIAAHSERLIGIALGAGD VRN.RTERSPEGTELLFARCSII QAARSAGIQAFDTVYSDANNE AGFLQEAAHIKQLGFDGKSLI PRQIDLLHNLYAPTOKEVDHA RVVEAAEAAAREGLGVVSLN	24380	54748	A	24518	1	1954	RQLNGVQLAFRESFHTVTRVTE QHAAGAVAIHQHRNQLLTGAL GIFAVAVCCLQQRLDILLADHL AEHVQFVIAQFVTCQQHGNGV RNRTIIFLLFNELRKIVETVRVQ QTQTGEVAFHPQLFRGRVVYG PDLNARRGNGVINCSTLEDPQF
MLRVIFRVNVETEAQQHESSR NGGALIKTINIRVENPILHDLA KDVPILNGDKIICSRLKREETLI VPSPDTIIQLGDLLHLDSRGSPI DTQDIDLQINSSVEKQFGDAIR TILDVLARYNVRGVQLNVDDD GALDCILRARLEALLARASGLI MISASLQQRKTRTRRSMLFVP ANAAMVNNSFIYPADALMFDI EDSVALREKDTARRMVYHALL HPLYRDIETIVRVNALDSEWG NDLEAVVRGGADVVRLTKTD AQDVLDIEKEILRIEKAGGREF GSTGLLAAIESPLGITRAVEIAH ASERLIGIALGAEDYVRNLRTE RSPEGTELLSARCPIWQAARSA GIQAFDTVYSDANNEAGFL/QI 24381 54749 A 24519 5121 5492 24382 54750 A 24520 2520 3525 24383 54751 A 24521 I 1263 MAINSPLNIAAQPGKTRLRKSI KLWQVVMMGLAYLTPMTVFI TFGIVSGISDGHVPASYLLALA VLFTAISYGKLVROFPEAGSAN TYAQKSINPHVGFMVGWSSLL DCLLAVVIIAAIYVGFSLAGFH YIGMLFVPGANAAMVSNSFIY ADALMFDLEDSVALREKDTAAI RMVYHALQHPLYRDIETIVRV ALDSEWGWNDLEAVVRGGAD VVRLPKTDTAQDVLDIEKEILR EKACGREPGSTGLLAAIESPLG TRAVEIAHASERLIGIALGAED VRNLRTERSPEGTELLFARCSII QAARSAGIQAFDTVYSDANNE AGFLQEAAHIKQLGFDGKSLIN PROJDLLHNLYAPTQKEVDHA RVVEAAEAAAREGLGVVSLN							SGLRLNLFAVLIVIIGGLVTAIL HKLFDIPLPVVLGIFSGAVTNTP ALGAGQQILRDLGTPMEMVDQ
TILDVLARYNVRGVQLNVDDI GALDCILRARLEALLARASGLI MISASLQQRKTRTRSMLFVP ANAMWNNSFIYPADALMFDI EDSVALREKDTARRMVYHALI HPLYRDIETIVRVNALDSEWG¹ NDLEAVVRGGADVVRLTKTD AQDVLDIEKEILRIEKACGREF GSTGLLAIESPLGITRAVEIAF ASERLIGIALGAEDYVRNLTTE RSPECTELLSARCPIWQAARSA GIQAFDTVYSDANNEAGFL/QI 24381 54749 A 24519 5121 5492 24382 54750 A 24520 2520 3525 24383 54751 A 24521 I 1263 MAINSPLNIAAQPGKTRLRKSI KLWQVVMMGLAYLTPMTVFI TFGIVSGISDGHVPASYLLALA VLFTAISYGKLVRQFPEAGSAY TYAQKSINPHVGFMVGWSSLL DCLLAVVIIAAIYVGFSLAGFH YIGMLFVPGANAAMVSNSFIY ADALMFDLEDSVALREKDTAI RMVYHALQHPLYRDIETIVRV ALDSEWGVNDLEAVVRGGAD VVRLPKTDTAQDVLDIEKEILE EKACGREPGSTGLLAAIESPLG TRAVEIAHASERLIGIALGAED VRNLRTERSPEGTELLFARCSII QAARSAGIQAFDTVYSDANNE AGFLQEAAHIKQLGFDGKSLIN PRQIDLLIHLLYAPTQKEVDHA RVVEAAEAAAREGULGVVSLN							MLRVIFRVNVETEAQQHESSRT NGGALIKTINIRVENPNLHDLAI KDVPILNGDKIICSRLKREETLK VPSPDTIIQLGDLLHLDSRGSPL
EDSVALREKDTARRMVYHALI HPLYRDIETIVRVNALDSEWG NDLEAVVRGGADVVRLTKTD AQDVLDIEKIELRIEKACGREF GSTGLLAAIESPLGITRA VEIAH ASERLIGIALGAEDYVRNLRTE RSPEGTELLSARCPIWQAARSA GIQAFDTVYSDANNEAGFL/QI 24381 54749 A 24519 5121 5492 24382 54750 A 24520 2520 3525 24383 54751 A 24521 I 1263 MAINSPLNIAAQPGKTRLRKSI KLWQVVMMGLAYLTPMTVFI TFGIVSGISDGHVPASYLLALA VLFTAISYGKLVRQFPEAGSAY TYAQKSINPHVGFMVGWSSLL DCLLAVVIIAAIYYGFSLAGFH YIGMLFVPGANAAMVSNSFIY ADALMFDLEDSVALREKDTAI RMVYHALQHPLYRDIETIVRV ALDSEWGVNDLEAVVRGGAD VVRLPKTDTAQDVLDIEKEILF EKACGREPGSTGLLAAIESPLG TRAVEIAHASERLIGIALGAED VRNLRTERSPEGTELLFARCSII QAARSAGIQAFDTVYSDANNE AGFLQEAAHIKQLGFDGKSLIN PRQIDLLHNLYAPTQKEVDHA RVVEAAEAAAREGVLGVVSLN							TILDVLARYNVRGVQLNVDDK GALDCILRARLEALLARASGLP MISASLQQRKTRTRRSMLFVPG
GSTGLLAAIESPLGITRAVEIAH ASERLIGIALGAEDYVRNLRTE RSPEGTELLSARCPIWQAARSA GIQAFDTVYSDANNEAGFL/QE 24381 54749 A 24519 5121 5492 24382 54750 A 24520 2520 3525 24383 54751 A 24521 I 1263 MAINSPLNIAAQPGKTRLRKSL KLWQVVMMGLAYLTPMTVFI TFGIVSGISDGHVPASYLLALA VLFTAISYGKLVRQFPEAGSAY TYAQKSINPHVGFMVGWSSLL DCLLAVVIIAAIYVGFSLAGFH YIGMLFVPGANAAMVSNSFIY ADALMFDLEDSVALREKDTAH RMVYHALQHPLYRDIETIVRV ALDSEWGVNDLEAVVRGGAD VVRLPKTDTAQDVLDIEKEILR EKACGREPPGSTGLLAAIESPLG TRAVEIAHASERLIGIALGAED VRNLRTERSPEGTELLFARCSII QAARSAGIQAFDTVYSDANNE AGFLQEAAHIKQLGFDGKSLN PRQIDLLHNLYAPTQKEVDHA RVVEAAEAAAREG\LGVVSLN							ANAAMVNNSFIYPADALMFDL EDSVALREKDTARRMVYHALQ HPLYRDIETIVRVNALDSEWGV NDLEAVVRGGADVVRLTKTDT
24381 54749							AQDVLDIEK\EILRIEKACGREP GSTGLLAAIESPLGITRAVEIAH ASERLIGIALGAEDYVRNLRTE RSPEGTELLSARCPIWQAARSA GIOAEDTYYSDANNEAGEL/OE
24382 54750 A 24520 2520 3525 24383 54751 A 24521 I 1263 MAINSPLNIAAQPGKTRLRKSI KLWQVVMMGLAYLTPMTVFI TFGIVSGISDGHVPASYLLALA VLFTAISYGKLVRQFPEAGSAY TYAQKSINPHVGFMVGWSSLL DCLLAVVIIAAIYVGFSLAGFH YIGMLFVPGANAAMVSNSFIY ADALMFDLEDSVALREKDTAFI RMVYHALQHPLYRDIETIVRV ALDSEWGVNDLEAVVRGGAD VVRLPKTDTAQDVLDIEKEILR EKACGREPGSTGLLAAIESPLG TRAVEIAHASERLIGIALGAED VRNLRTERSPEGTELLFARCSII QAARSAGIQAFDTVYSDANNE AGFLQEAAHIKQLGFDGKSLIN PRQIDLLHNLYAPTQKEVDHA RVVEAAEAAAREG\LGVVSLN	24381	54749	A	24519	5121	5492	GIQNI DIVIODILINOI DI QE
KLWQVVMMGLAYLTPMTVFI TFGIVSGISDGHVPASYLLALA VLFTAISYGKLVRQFPEAGSAY TYAQKSINPHVGFMVGWSSLL DCLLAVVIIAAIYVGFSLAGFH YIGMLFVPGANAAMVSNSFIY ADALMFDLEDSVALREKDTAF RMVYHALQHPLYRDIETIVRV ALDSEWGVNDLEAVVRGGAD VVRLPKTDTAQDVLDIEKEILR EKACGREPGSTGLLAAIESPLG TRAVEIAHASERLIGIALGAED VRNLRTERSPEGTELLFARCSII QAARSAGIQAFDTVYSDANNE AGFLQEAAHIKQLGFDGKSLIN PRQIDLLHNLYAPTQKEVDHA RVVEAAEAAAREGYLGVVSLN			-			3525	
24384 54752 B 24522 62 405 GKMVDGPVIDRARLVLSRAEL	24383	54751	Α	24521		1263	MAINSPLNIAAQPGKTRLRKSL KLWQVVMMGLAYLTPMTVFD TFGIVSGISDGHVPASYLLALAG VLFTAISYGKLVRQFPEAGSAY TYAQKSINPHVGFMVGWSSLL DCLLAVVIIAAIYVGFSLAGFHV YIGMLFVPGANAAMVSNSFIYP ADALMFDLEDSVALREKDTAR RMVYHALQHPLYRDIETIVRVN ALDSEWGVNDLEAVVRGGAD VVRLPKTDTAQDVLDIEKEILRI EKACGREPGSTGLLAAIESPLGI TRAVEIAHASERLIGIALGAEDY VRNLRTERSPEGTELLFARCSIL QAARSAGIQAFDTVYSDANNE AGFLQEAAHIKQLGFDGKSLIN PRQIDLLHNLYAPTQKEVDHAR RVVEAAEAAAREG\LGVVSLN GKMVDGPVIDRARLVLSRAELS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24385	54753	I _A	24523	891	3240	RYRKSPGSWR/CMSCILTSWITT
24363	34733	^	24323	0.71	3240	MFPI*RLGQWC*NPCVTMAL/L
						APIDVHLMVKPVDRIVPDFAAA
						GASIITFHPEASEHVDRTLQLIK
						ENGCKAGLVFNPATPLSYLDY
						VMDKLDVILLMSVNPGFGGQS
						FIPQTLDKLREVRRRIDESGFDI
		1				RLEVDGGVKVNNIGEIAAAGA
						DMFVAGSAIFDQPDYKKVIDE
						MRSELAKFEDIRGVAFDLDGTL
						VDSAPGLAAAVDMALYALELP
						VAGEERVITWIGNGADVLMER
						ALTWARQERATQRKTMGKPPV
		i				DDDIPAEEQVRILRKLFDRYYG
				1		EVAEEGTFLFPHVADTLGALQA
						KGLPLGLVTNKPTPFVAPLLEA
:						LDIAKYFSVVIGGDDVQNKKPH
						PDPLLLVAERMGIAPQQMLFVG
						DSRNDIQAAKAAGLDAQARRK
						NAGYYYYLRLAKRQMRAGVA
						EALRDDLLAAKLNVKLVNAGD
						YKFKQIASEKLLIVVTSTQGEGE
						PPEEAVALHKFLFSKKAPKLEN
						TAFAVFSLGDSSYEFFCQSGKD
						FDSKLAELGGERLLDRVDADV
						EYQAAASEWRARVVDALKSRA
						PVAAPSQSVATGAVNEIHTSPY
						SKDAPLVAWRLPNVEGRARAG
						GASSFLADRVEEEGEVRVFIEH
						NDNFRLPANPETPVIMIGPGTGI
						APFRAFMQQRAADEAPGKNWL
						FFGNPHFTEDFLYQVEWQRYV
24386	54754	Α	24524	950	2124	MSLISSVIAATWLHVKSYRLSC
						GFYACQSLVLVS/IFPTLSCLFA
		1				AEQLLIWSASAFITKVLLVPLIM
						TYAARNIPQNIPEKALFGPAMM
						ALLAALIVLLCAFVVQPVKLPM
						ATGLKPALAVALGHFLLGLLCI
1						VSQRNILRQIFGYCLMENGSHL
						VLALLAWRAPELVEIGIATDAIF
			8			AVIVMVLLARKIWRTHGTLDL
						TPLLFSLLCFACRKRRLSATRTV
	}					TVLHSLGITLLLILALWVVQTA
						ADAGEIFAAGLWLHIDGLGGLF
	1					LAILGVIGFLTGIYSIGYMRHEV
	ļ 					AHGELSPVTLCDYYGFFHLFLF
		1				TMLLVVTSNNLIVMWAAIEAT
						TLSSAFLVGIYGQRSSLEAAWK
						YIIICTVRVAFGLFGTVLVYANA
1						ASVMPQAEMAIFWSE

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	<u> </u>	<u> </u>	<u> </u>			
24387	54755	Α	24525	1	757	RWRLPGVPPFNGFFSKFPLFAP
						GFALSVEYWILLPAMILLMIESV
						ASFAWFIRWFGRVVPGKPSEAV
						ADAAPLPGSMRLVLIVLIVMSLI
						SSVIAATWLHVKSYRLSCGFYA
						CQSLVLVSIFATLSCLFAAEQLL
		1				IWSASAFITKVLLVPLIMTYAAR
						NIPQN\IPEKALFGPAMMALLA
						-
						ALIVLLCAFVVQPVKLPMATGL
	÷					KPAL/AGLATDAIFAVIVMVLL
	1		ŀ			ARH\IWRTHGTLDVNNLHALK
		<u> </u>				GIMSCSYSS
24388	54756	A	24526	249	3943	
24389	54757	A	24527	1	1278	
24390	54758	В	24528	48	144	
24391	54759	В	24529	27	239	
24392	54760	Α	24530	1	551	
24393	54761	Α	24531	3	258	MLFDEPTSALYPEMIN*VLDVM
						VELANEGMTMMVVTHEMGFA
						RKEANRVIFMDEGKIVEYSPKD
						AFFDDPKSDRAKDFLAKILH
24394	54762	Α	24532	2	979	
24395	54763	A	24533	Ī ₁	4116	
24396	54764	A	24534	li -	1195	MLIAEGLMTAKNITVITNSLPA
21370	13 77 0 1	` `	2.55.		1133	AFALSENKDITLVVCGGTVRHK
						TRSMHGSIAERSLQDINADLMF
	1	1				
						VGADGIDAVNGITTFNEGYSIS
						GAMVTAANKVIAVLDSSKFNR
						RGFNQVLPIEKIDIIITDDAVSEV
		1				DKLALQKTRDTSLVYVLSLADF
						FRTASTIGERDGTQVEMILFAGF
		1				VYLVISLSASLLVLTDCSTEVK
		1				KGEVVVVCGPSGSGKSTLIKTV
		1				NGLEPVQQGEITVDGIVVNDKK
						TDLAKLRSRVGMVFQHFELFPH
		1				LSIIENLTLAQVKVLKRDKAPA
		l				REKALKLLER/VSGFFAHANKF\
		İ				PAQLSGGQQQRVAIARALCMD
	į.					PIAMLFDEPTSALDPEMINEVLD
						VMVELANEGMTMMVVTHEM
	1					GFARKVANRKHTTRSTIALQSL
0.400=	64765	 	04505	202	0.62	ATCT
24397	54765	A	24535	383	962	TDNDRPGLGTDMKKASAIVTN
·						VAVVLSGRYRRERASGEWAAL
		1				SFLRPHTLTVFVIPHIQWDDPQ
		1				MSRP\SGRHLFRCRTVQRKHAL
						QVVEKFRPFFPSDRYRYGAPGD
						YAQSFGGQIDDVVKAIQTPSKF
						RRFLRVSFD*IRLPILHSGKSPLR
						RMPNLMRRHSHTARHYPALGT
		1				RGPQPTDRQERPAVASGQ
L		İ		I		NOI QUIDIQUIA TABOQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24398	54766	A	24536	1477	2387	
24399	54767	Α	24537	1	1560	
24400	54768	A	24538	305	718	
24401	54769	A	24539		2161	MNDVDRVGGKNASLGEMITNL SGMGVSVPNGFATTADAFNQF LDQSGVNQRIYELLDKTDIDDV TQLAKAGAQIRQWIIDTPFQPEL ENAIREAYAQLSADDENASFAV RSSATAEDMPDASFAACTLVPG QNLSTSNKDVIELPDNQYDLDK MVNIYPVTPGLIDQLRAKPIMS QANPELEQQIANYEYRIGIGDV LMVTVWDHPELTTPAGQYRSA SDTGNWVNADGAIFYPYIGRLK VAGKTLTQVRNEITARLDSVIE SPQVDVSVAAFR\SQKAYV\TA EVSKSGQQPITNIPLTIMDAINA AGGLTADADWRNVVLTQNGV KTKVNLYALMQRGDLRQNKLL HPGDILFIPRNDDLKVFVMGEV GKQSTLKMDRSGMTLAEALGN AEGMNQDVADATGIFVIRATQ NKQNGKIANIYQLNAKDASAM ILGTEFQLEPYDIVYVTTAPLAR WNRVISLLKKLERLTSIAITLPGI IDTENGIVHRMPFYEDVKEMPL GEALEQHTGVPGEAKGLVGAP PGARDVIQVVSIPNVGRAVIPN GHLLPQGSSSLVEIAHTQGGPL
						MGKRCYCGNHGCLETIASVDSI LELAQLRLNQSMSSMLHGQPL TVDSLCQAALRGDLLAKDIITG VGAHVGRILAIMVNLFNPQKILI GSPLSKAADILFPVISDSIRQQA LPAYSQHISVESTQFSNQGTMA
24402	54770	С	24540	1	1389	
24403	54771	A	24541	405	576	CEMGAPPWPGCWLS\LFDVDG RFVLAWMLFQGSYLAGALGDQ PVHIHLGQLLALWRPL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24404	54772	A	24542		1127	MARTLAHELIDFGEIKRGLLGIK GTEMSADIAKAFNLDVQRGAF VSEVLPGSGSAKAGVKAGDIIT SLNGKPLNSFAELRSRIATTEPG TKVKLGLLRNGKPLEVEVTLDT STSSSASAEMITPALEGATLSDG QLKDGGKGIKIDEVVKGSPAAQ AGLQKDDVIIGVNRDRVNSIAE MRKVLAAKPAIIALQIFPVDHR GAIGGQTTDIPPDPSGINHDRC MKGLSFAGEMVTYTNTAHRRA
						FTNQPRDFMSVAWLGKGIAILP SVGEVRSPVAGRIASLFATLHAI GIESDDGVEILIHVGIDTVKLDG QIFSAHVNGEAGLKRVVKKEH AD/VQRHAESV*RRGQAQGTD GIRRQDNRVQPGCERCTHQLLR
24405	54773	A	24543	95	599	QIMRMLNGSSTPNTSLPTPQAF DRSLPINATSARPSSTSTRHSAE SSISRDCVSNVS/HSGG*TGGIK G*CHAHPADLTACHEADSRGSP SHAVAAVALHVID/GWYLGDQ TAAYATGSAVSCPSSGNQFSRP SAAGRFSAVSAPQGAQRHPGR ARRRPRRRRTAGGCRR
24406	54774	A	24544	640	890	LANAQLGSNEISIFQCFAAIQRFI KLHRDASVINHPLAE/TRK*YPV SVYHLRHPPAIIR*PAIRDNA**I LPVTPGYSCSRHQ
24407	54775	A	24545	3	257	ETDIFICDSREFFHHFAKASVHI* PVVSIANFAIQSGKILALGAENI ATTSDRRAQRQKAVCKAHDAS SVSFRTLSVRPTNDCP
24408	54776	A	24546	332	1047	RTRPYCPGMGNEPMTGWMAA AVVTLMIRMCFSVYTMLSESC QRMVIVGYGRCFADRQNLMVC LRSMPNVFTGSCARMRCCLSEN LLYRHRNGHIQAEWP*KKAISD VKQYRTSCWERWNAASATIFR/ LSPVEWLTDNGSCYRANETRQ FARMLGLEPKNTAVRSPESNGI AESFVKTIKRDYISIMPKPDGLT AAKNLAEAFEHYNEWHPHSAL GYRSPREYLRQRACNGLSDNR CLEI
24409	54777	A	24547	1200	1544	LTGVADGNVDKPCYKFARLW YWIYPAFPPIRRVKVAYREYRP PTTPRVRQVRDLTFCHRPAIPDR MHRY*SSHRALPGCWSKVPAQ KSSRTVSGTD*VDLLSPRAQRA PSPLQG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24410	54778	A	24548	263	625	
24411	54779	A	24549	742	851	VSAQQRVILNKIAVLNRK*RYG CWRLWICPYISRHLLSLNPLQA RCRRYSRGER
24412	54780	A	24550	1	532	MKTLLIIDANLGTRLYGEDPAG AAARKAKLEIIDNPNDAEMAIV LGDSIPNDSALNGKNVWLGDIS RAVAHPELFLSERKGHAKPYTA PVAATAPVGASGPKRVVAVTA CPTGVAHTFMALKPLKPKRKN VAGIPDTGQYCETVQTSCW/AA VETPLRQRSSVVSSGVADG*WF MLPG
24413	54781	В	24551	1	888	
24414	54782	Α	24552	1	2283	
24415	54783	A	24553	992	1662	KKAISDGALTGSSSAVITERDC VSRSRWTAVIVRHCTGRSLPAA STVKQYRTSCWERWNAASATI FR/LSPVEWLTDNGSCYRANET RQFARMLGLEPKNTAVRSPESN GIAESFVKTIKRDYISIMPKPDG LTAAKNLAEAFEHYNEWHPHS ALGYRSPREYLRHGLVSFISSLQ KFINDFAFEDGLFSGYDAEKRQ YEKSSWNYQLDENGYAKRDET LTHPRC
24416	54784	Α	24554	1	2262	
24417	54785	Α	24555	158	410	
24418	54786	В	24556	1	2607	
24419	54787	A	24557	715	1016	KSAAVPRSGASSQSDSAHFAQP EYAAFSRFPRGSFRHPRAA*YF GRLNHHVRSERGINP*YHRFNL HWRYVADAISADGAFRTVSGG GGRRSGAGRVDWQ
24420	54788	A	24558	652	1214	RTDHCVSRSRWTAVIVRHCTG RSLPAASTVKQYRTSCWDRWN AASATIFR/LSPVEWLTDNGSCY RANETRQFARMLGLEPKNTAV RSPESNGIAESFVKTIKRDYISIM PKPDGLTAAKNLAEAFEHYNE WHPPSALGYRSPREYLGQRACT GHAIGSGPSSAFFGAPVRRVSRT SVTAKACPRGLL
24421	54789	Α	24559	647	1976	
24422	54790	A	24560	289	574	ETPASSPTLRTSIPSISWMTRGSL RRVMLLCPFPSESGTVSEKAWP EWSSFSSSPPSCRTSASSPPSHLR /NIDVSPKHVVFATIPRNYTMSF LPR
24423	54791	Α	24561	816	1375	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24424	54792	Α	24562	1	3993	MSQKLKVVTIGGGSSYTPELLE
						GFIKRYHELPVSELWLVDVEGG
						KPKLDIIFDLCQRMIDNAGVPM
						KLYKTLDRREALKDADFVTTQ
						LRVGQLPARELDERIPLSHGYL
						GQETNGAGGLFKGLRTIPVIFDI
						VKDVEELCPNAWVINFTNPAG
						MVTEAVYRHTGFKRFIGVCNIS
			i			IGMKILKTRTLDTTGSTSTGDTL
			}			LTQSLLMELSALCRVEVEEGLA
		1				LVALIGNDLSKACGVGKEVFG
		l				VLEPFNIRMICYGA
24425	54793	Α	24563	1136	3608	KKAISDGALTGSSSAVITERDC
						VSRSRWTAVIVRHCTGRSLPAA
						STVKQYRTSCWERWNAASATI
	1					FR/LSPVEWLTDNGSCYRANET
						RQFARMLGLEPKNTAVRSPESN
	}					GIAESFVKTIKRDYISIMPKPDG
						LTAAKNLAEAFEHYNEWHPHS
						ALGYRSPREYLRQRACNGLSD
		ļ				NRCLEIYGQIQLGTGELNITTGV
						VVTARDTHIALNFQKRRTTGLA
						TQSRQRCQNRRRVMGKIVIYRD
						AIRFTTQLQTATSINKRTQRVSG
						ICRRYTNMASRCNRHQAIVHIM
						FAYQRPFHFAHFFAIQQNFPFGS
						IGGQLFRLPVTLLTHQLLLAPA
						AHRHGLFQVDVIFRPDNTALPR
						NDTHQMVELFLDRFQVVKDIG
						VIELKVVEDQRTRAVMNKFRA
	İ					FVEKGAVILIRFDKKKDIFLQYR
						FHTRISTGHGVANNHQIRLRLE
			•			LAGIVPLNQLNPLPLNSVPMINI
						MNGGENADNNVDIQEFMIQPV
						GAKTVKEAIRMGSEVFHHLAK
						VLKAKGMNTAVGDEGGYAPN
		1				LGSNAEALAVIAEAVKAAGYE
						LGKDITLAMDCAASEFYKDGK
		1				YVLAGEGNKAFTSEEFTHFLEE
		1				LTKQYPIVSIEDGLDESDWDGF
	1					AYQTKVLGDKIQLVGDDLFVT
	1					NTKILKEGIEKGIANSILIKFKAQ
						HAGLLHMYTCAMLGVGGQCW
	1		L			RYCMSQAFKEDRFAQDQRVAV
24426	54794	В	24564	83	2851	
24427	54795	В	24565	1	1578	
24428	54796	Α	24566	574	753	
24429	54797	Α	24567	1	2112	
24430	54798	В	24568	27	969	
24431	54799	Α	24569	1	1773	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24432	54800	A	24570	822	984	PSALCGCHSL*CSNASARFFAA VNPSGAFGCRADHPDGELFLTG YVADGGYHS
24433	54801	Α	24571	1	4287	
24434	54802	Α	24572	1	2384	
24435	54803	A	24573	3	3682	ACSARPSEAELLELIDTLNADN TIDGILVQLPLPAGIDNVKVLER IHPDKDVDGFHPYNVGRLCQR APRLRPCTPRGIVTLLERYNIDT FGLNAVVIGASNIVGRPMSMEL LLAGCTTTVTHRFTKNLRHHVE NADLLIVAVGKPGFIPGDWIKE GAIVIDVGINRLENGKVVGDVV FEDAAKRASYITPVPGGVGPMT VATLIENTLQAMSNMATFSLGK HPHVELCDLLKLEGWSESGAQ
24436	54804	A	24574	1	1382	AKIAIAEGQVKVD MLRGVNVLADAVKVTLGPKG RNVVLDKSFGAPTITKDGVSVA REIELEDKFENMGAQMVKEVA SKANDAAGDGTTTATVLAQAII TEGLKAVAAGMNPMDLKRAS RSRKGYGPARIRQELNQKGISR EATEKAMRECDIDWCA**RF*S DKHHHRT*RQ/PTRHRYPDESV PGSHNHRSAHALSSAQGYILLP RAVLSGWRFENVHPTPGYPAG SGSVFVPPPMTSRRLPAPDNFA APHRYRVLKTGLLSPGCHVRN
						ADTL
24437	54805	A	24575	1104	1215	LAIPMKDFITEAWLRANHTLSE GAEIHLPADSRLTPSARELLESR HLRIKFIDEQGRLFVDDEQQQP QPVHGLTSSDEHPQACCELCRQ PVAKK\PDTLTHLSAEKMVAKS DPRLGFRAVLDSTIALAVWLQI ELAEPWQPWLADIRSRLGNIMR ADALGEPLGCQAIVGLSDEDLH RLSHQPLRYLDHDHLVPEASHG RDAALLNLLRTKVRET\ETVAA Q\VFITRSF\EVLRP\DILQALNRL SSTVYVMMILSVTKQPLTVKQI QQRLGETQ
24439	54807	A	24577	243	644	HAGSVCPSLYRCQRTAKRSGFY LLLKRLWSFSGNI*IHSGQQNPG RNQQLGLPRSRKGESSRAHPSC QRRPQVWPHPLAGSRPPRPWA ACARLGPIAPARQIPERWPGQV PVLADAHVLPCLAGSLRPALYD F

SEQ ID		1	SEQ ID NO:		1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24440	54808	A	24578	237	454	RSRLPEGVPVCWCNHSVR*RDC RFPVGKLHRLWL/LALPGVRLIF RASTKRITGYINARSASIASASA RNRLV
24441	54809	A	24579	377	1155	KAYSFATMKSLAISCRWQNA*S HVSTFVMVRW/LKGVQFRNHEI IGDIVPLAKRYAEEGADELVFY DITASSDGRVYPTLITRLADRFG VQCIVVGIDTWYDARNLKDISV VGLLGAQVPSVRAGMPQSRRQ TEGAQGMVRNEEGGSLRLSHH QACKATHTQQWTLEVTAQSSY NLFPEQWVKLFWGIRQGDLPL AFSQKLSRQTFQQQQTQTPRPI VGQAPKDLQTGTGTPQTHAVH SGDHAAGEQSCKSCKAPIYPLT AQFSLETRSFWGSEWGYSTHG ELRRRDKGRRFY
24442	54810	A	24580	1107	1458	RTFWWTTCNSLAPCAGGYPST GAALPVSRKLPGRGLLLSGWG EHNARYRAGTFLWALFCAAAV AYWRRC\WGQYGMWFLRIRRG TLGDGTIRRVFVGSLWAAVVIV EIGVEHIGVFKA
24443	54811	A	24581	171	352	PPAPPAESDGLPAHTTRARTTA DDLRNKRRQHVGC*YRTRGSP RPNRTWWRVCRYRSPHR
24444	54812	A	24582	5	226	
24445	54813	A	24583	476	1939	RPESQHNTAKPAMKVLEQARR FLGIRWLSAPASLAN/PVILLVV GNILNIVLDVWLVMGLHMNVQ GAALATVIAEYATLLIGLLMVR KILKLRGISGEMLKTAWRGNFR RLLALNRDIMLRSLLLQLCFGAI TVLGARLGSDIIAVNAVLMTLL TFTAYALDGFAYAVEAHSGQA YGARDGSQLLDVWRAACRQSG IVALLFSVVYLLAGEHIIALLTS LTQIQQLADRYLIWQVILPVVG VWCYLLDGMFIGATRATEMRN SMAVAAAGFALTLLTLPWLGN HALWLALTVFLALRGLSLAAI WRRHWRNGRPGTGCLNTKGS QQQHRHDSTTQHIHLQLLVIRV KTTTTNNSQQTPQYPCQQYHT DSRQHRQQNRMVCQRIPHALPS PRFRQISRGSTAKVPCAKARAF LRVKSDPTLITRLADRFGVQCIV VGIDTWYDAETGKYHVNQYTG DESRTRVTQWETLDWVQEVQK RGAGEIVLAYVSE

SEQ ID NO:	SEQ ID NO: of peptide sequence	į.	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24446	54814	Α	24584	1	50	
24447	54815	A	24585	354	456	HLFSEACCSPGVLRELCRPF*SP GTDRNLVMHGH
24448	54816	A	24586	494	835	ITWLPGFRPPFQGSEQFCLAGVP GATGGSESVLHTRNSYKMLEV RREVAALISALKVSAKCRNCGS SHALTEP/ASSPCWRETSTRPTG APAPTIPPPAASPSPGPMCVQPH GED
24449	54817	Α	24587	3	726	
24450	54818	Α	24588	1	1635	
24451 24452	54819	A	24589 24590	362	517	RTTHAPASTPNTGTMLCRPYPC CWRRKTP\IIFRATPQWFVSMD QKGLRAQSLKEIKGVQWIPDW GQARIESMVANRPDWCISRQRT WGVPMSLFVHKDTEELHPRTL ELMEEVAKRVEVDGIQAWWD LDAKEILGDEADQYVKVPDTL DVWFDSGSTHSSVVDVRPEFA GHAADMYLEGSDQHRGWFMS SLMISTAMKGKAPYRQVLTHG FTVDGQGRKMSKSIGNTVSPQD VMNKLGADILRLWVASNDYTG EIAGSDENLKRACHYNACSISL ARRH TKSLISSMSSISSPPVVLICRNSS PEV**RKI*LGTELEFVIMVIAVS
						IAWP
24453	54821	A	24591	250	930	
24454	54822	A	24592	1388	1912	VCPAAMTGFAGSPFCTAVVTFP DITPLLWILMC*LAILMAESEPK AISPPTISVVP*YAAESLLRVRIP VPSFVTSPLPLILCVIAQLRSVE MVSRPLLITEASPRFFCSLTLSA LPVSMLNCPTKRLLSPSNSTSVL AVALTIPFPLMLLRSSPVTFFSV NRPSWAIFPLPNP
24455	54823	В	24593	1	6018	
24456	54824	A	24594	510	1003	VALQADQKHKHERPLPPPIPTM FQLAQPGVQLHAKPVYQVP/V MMQATEPTGI*VRIPVLSISTNA PSITSIWPLFVRKISVLTLASVV AASHAASSPRQCAIRCKIWGRW AGSLRRDDGFGVRSRGSRYGA SVSISNFSAGINGKSSRSSSPRRS SQTQPVMPI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24457	54825	A	24595	197	4314	RASTELSLRCCAPSLQ/ALLHFL
						YLACFTWMLLEGLHLFLTVRN
		ŀ				LKVANYTSTGRFKKRFMYPVG
						YGIPAVIIAVSAIVGPQNYGTFT
					:	HCWLKLDKGFIWSFMGPVAVII
						LINLVFYFQVLWILRSKLSSLNK
						EVSTIQDTRVMTFKAISQLFILG
						CSWGLGFFMVEEVGKTIGSIIA
						YSFTIINTLQGVLLFVVHCLLNR
						QINSLLLTWTLWILRQRLSSVN
						AEVSTLKDTRLLTFKAFAQLFIL
Ì						GCSWVLGIF
24458	54826	Α	24596	1409	1573	
24459	54827	Α	24597	1283	1563	
24460	54828	Α	24598	281	839	CVPTTDLVVQSQI*PRWRNWPR
		ŀ				CSCSHVSMKAASSPERTFTYCS
						GIYWLISGSGIGVSVPLLISIGES
						VINLAVRPVIAHAIRVAITTRQT
			-			TRVAVTVVVALPVFRAVEHHV
						FQHLTQTRTGFRWHQFFFNRRQ
						REQSFTQGEGQRWANFQNGKV
						VNHQLAQRLRRLIIGRVRFENQ
						RFVVFDHLA
24461	54829	В	24599	1	5652	
24462	54830	A	24600	559	1665	RVERKATIEMKSQRTTSDSWLII
						LAKQTGIIRAAKKGKKTFILHD
						GPPYANGSIHIGHSVNKILKDII
		İ				VKSKGLSGYDSPYVPGWDCHG
						LPIELKVEQEYGKPGEKFTAAE
						FRAKCREYAATQVDGQRKDFI
						RLGVLGDWSHPYLTMDFKTEA
						NIIRALGKIIGNGHLHKGAKPV
						HWCVDCRSALAEAEVEYY\DK
	1					TSPSIDV/AHPFMGFDVPAILGD
	İ					HVTLDAGTGAVHTAPGHGPDD
						YVIGQKYGLETANPVGPDGTY
						LPGTYPTLDGVNVFKANDIVVA
		1				LLQEKGALLHVEKMQHSYPCC
1						WRHKTPIIFRATPQWFVSMDQK
] .					GCDIGFHASSNGKRRADDSTDN
	1					HVRRLCARQRGQPHPHQFQRT

3231

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24463	54831	Α	24601	1	1190	MTDFRESDTMSLFDKKHLVSP
						ADALPGRNTPMPVATLHAVNG
						HSMTNVPDGMEIAIFAMGCFW
						GVERLFWQLPGVYSTAAGYTG
	!					GYTPNPTYREVCSGDTGHAEA
						VRIVYDPSVISYEQLLRKYRCV
						AVALLGYFKCKPILLNPSYKSM
						QDDLAFIARFHFKDLKFRRFSL
						KADQKSRIYERVLSIAGVNSWK
						DKQHQPRLVEHLLVCAESWVA
						PRALFDAAIEYLAHQKIAIPAYL
						VAFLVLGPLVLSLIPSQMLPSLT
						WSSITYSIVVIVEAGYQYHSRLR
						RFGRVEVNEQIEYLIHNPIRTRA
						RAVNFVDNNNRLQAPRILRQDS
						WWITRPLPGLTWAIIGSPGILPQ
						VVKSARVMKQAVAYLEPFIEA
						SKEQGKTNGKMVIATVKGDVH
						DIGKNIVGVVLQCNNYEIVDLG
						VMVPAEKILR*KLGGTPRTL*R
						SNRVPGPPEDRHSCLFGCV/YW
1						FLVP*CCLSSLRRCCRR*RGQAS
						RIQ*LLSSKLAISTIL/GLRRFGR
	!					VEVNEQIEYLIHNPIR/IARQGG
						QLC**QQSASGPTDITAGFMVD
						HQTFTGADVGDNRVARNPATG
						GQIGARHETGGGLPRTVY*SQQ
						RAGQNQRQDGDRHREGRRPRH

3232

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24464	54832	Α	24602	1	1591	MGILVSPLLAPLIGGLLDTMWN
						WRACYLFLLVLCAGVTFSMAR
						WMPETRPVDAPRTRLLTSYKTL
						FGNSGFNCYLLMLIGGLAGIAA
						FEACSGVLMGAVLGLSSMTVSI
						LFILPIPAAFFGAWFAGRPNKRF
						STLMWQSVICCLLAGLLMWIP
						DWFGVMNVWTLLVPAALFFFG
						AGMLFPLATSGAMEPFPFLAGT
						AGALVGGLQNIGSGVLASLSA
						MLPQTGQGSLGLLMTLMGLLI
		İ				VLCWLPLARGYRLSSWTSTTAS
						EDIDCEEGDGEGNETHGLQLAV
						EPEGLQPNYMGRGKIRFRDTCL
						LFCTQSRGSQWGFARTELVALL
		İ				PPFLSGAADYRAGLALWLARG
						ADCDVDERHRADRQSNLARSS
				-		GGFIALAAGAKSDRRLRELNQS
						LQKELARNQHLAERLLETEESV
						RRDVARELHDDIGQTITAIRTQ
						AGIVQRLAADNASVKQSGQLIE
						QLSLGVYDAVRRLLGRLRPRQ
						LDDLTLEQAIRSLMREMELEGR
						GIVSHLEW\QSMNQR*AKTSA*
						RCFVSARKG
24465	54833	Α	24603	1	1752	
24466	54834	Α	24604	1	699	
24467	54835	Α	24605	965	5516	
24468	54836	Α	24606	84	3760	
24469	54837	Α	24607	27	278	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24470	54838	A	24608	311	2529	REWGSTRRCERRISLSRNIARW RTIWRITRPCRRCVFRFIASTLR LHLHLIMNDLPLEEYPCVRDLV PGRFLKFL/HRDPKGTTLVLASD LPGRIPVDPRSIIVMLFLAVRWL QRQLAGQELLETRATRILNGER GSNVLGTIYEWPPRTSSALDTL LREIQNAREQHSRLDTLIRSYA AQDVKTGLNNRLFFDNQLATL LEDQEKVGTHGIVMMIRLPDFN MLSDTWGHSQVEEQFFTLTNL LSTFMMRYPGALLARYHRSDF AALLPHRTLKEAESIAVFRALS RLPDLWHRSIGTLTEALPQTDK RQEQIAVALTSYSGLYGLGYGS KEIQLTYSGTAGDLISPGLSAIH GRCILNTLVFIVKLRWMENTLH RFGTHHGCSAVRSSHLDSGSPR RSSWHMARARNHIHVLEENMA LLMEQAIANEGLFYRLLYLQRS LTAASSLDDMLMRFHRWARDL GLAELLVVLPEAKAVGSVAMS MLGSDADWGVVLFTSRDASHY QQGQGTQLLHEIALMLPELLER WIERGWEQQVWRYGFLRYVAF LTGWSAQNELKANPAKGVSAP KAPRHLPKNIDVDDMNRLLDID INDPLAVRDLMYGAGLRLSELV GLDIKHLDLESGEVWVMGKGS KERRLPIGRNAVAWIEHWLDL RDLFGSERERRQHAGTNVLQT
						ADQRASRARQEGERLHRAAGR
24471 24472	54839	A	24609 24610	1	2911	MNIDAKILNKILANRIQQHIKKL IHHNQVSFIPGMQGWFNIHKSM KVIQHINRTKDKNYMIISIDAEK AFDKIQQPFMLKTLKKLVAGM MAVAAPDDPPPSVSSSKPSTV APPGRISPSGRALCTSLPSALSSS SKFSLLRIFQVPAKLGMVAVAA AAGVNAMLRKVAVAAASKPH VEIRQDGDQFYIKTSTTVRTTEI NFKVGEGFEEETVDGRKCRVR
24473	54841	A	24611	249	609	PQSHYSVPVSPLGAHGPLLLEE PVSPFAACGAPS QIEVCVRIKPKAGTVLAYS*RSR QERPPAPGMTSLSLAAVSSDNP VSGDEHIRLARASGRFGIGRVA QPKRIQRRSSTPRAATDLGNPG HRPCTVRKSALLVVSVCRRHSS RSLRFRSC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	,	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24474	54842	Α	24612	942	1121	
24475	54843	Α	24613	1010	1658	EISSSKRSRTVCKRRAPMFSVDS
						FTCHAASAMRLIPESVNSIFTPS
						VAISALYCTVIEAFGSVRIRSKS
						LVVSACSSTRIG\DDLAVPGSGQ
						TVSQSGTRQKR*TECGQF*PSR
						VWWKRCNLLPAAAGHAERLH
						EKHPDRKTRRVRLVWLAGPPT
						RLLGTMGAGTLVLGACCLPWP
						REGAHNVATAACGGWRARCD
	1					LRFRLSCKLGLSRQFFGWNA
24476	54844	Α	24614	1	1995	
24477	54845	Α	24615	1	2853	
24478	54846	Α	24616	1	1755	
24479	54847	Α	24617	2942	3454	SVPDWLMCTAFLSLSRINSPRL
		1				LKAELLLTSSIPVLLPSLLLPRFQ
		1				VPSLFATAVPVSAAPLIIAPSV*L
		1				LKTPPADIIRPVTLPLLLNTPAFC
						TILLMVPVLFTNELSAPAFMVIC
		ŀ				AVISPALLRIPPLTSTLFVITPAF
						LMTAPLSSFTKLLMMPPLVFSR
		ļ				RPLDTVI
24480	54848	Α	24618	1	2959	MKRYRPREILPWKVPPGNYTPK
						IFKIKGTGHNPMSAIENFDAHTP
						MMQQYLRLKAQHPEILLFYRM
						GDFYELFYDDAKRASQLLDISL
						TKRGASAGEPIPMAGIPYHAVE
						NYLAKLVNQGESVAICEQIGDP
						ATSKGCCRSGNRNGGRKIVAVS
						LKKVSGAAKPCGHFRSTRYRV
i						AYSTIINNKAGIWSAFEGVSAT
l						RPNPTPRASPAVRTLSRGTPEPR
						KISTDADNLAQGSLKERLPAWL
						WGKKENSKWEVLD
24481	54849	Α	24619	441	1425	
24482	54850	Α	24620	1	2574	
24483	54851	В	24621	1	1725	
24484	54852	Α	24622	160	390	
24485	54853	A	24623	892	1183	
24486	54854	Α	24624	1	2493	A ODDAY MORGOD AND CARACTERS
24487	54855	Α	24625	435	829	ASDVLVSEGCR/YRGASALTST
						SVICINTDKALRKIIRLPLNGLR
					1	KPLNAMTPLPGITWPLCITTEKE
1						DLSISDRLSTCIVKFSHPEPEMS
						VRKFVPPAILLCSVCLRFSCPTR
<u> </u>	5.1055	 	0.160.5	7.62	1062	HQKAPAKWILKSCSAWMR
24488	54856	A	24626	763	1863	
24489	54857	Α	24627	1423	1712	

SEQ ID	-	I	SEQ ID NO:	1		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24490	54858	A	24628	1	2792	MTYLSAQDHELIQVVQNGHLQ
						MCGKSETGNEASPVKCIVLQVL
		İ				RGLQYLHRNFIIHSQEQTIPDPSI
						SDQQETPLLAGDMAQRGRRGG
						FYYVTAYIYVDKKAPVHMRTA
ļ						AQGLITLCCQGFGSLLGYRLGG
						VMMEKMFAYQEPVNGLTFNW
						SGMWTFGAVMIAIIAVLFMIFF
İ					,	RESDNEITAIKVMEDSMSSATT
						LTTPQEQVDSLIMQIAEENGLE
						VLDQLSQLPEGASAVGESSVRS
Ī						QEDQLSRSFICLMAMM
24491	54859	C	24629	213	345	
24492	54860	A	24630	1	474	MHQTTAKTPHTSYPNNYEEHN
						RRANRSKNYDRIRHTTIILKRTN
						PAQTITNLLKSRFNRYNYVQPR
						SCRSIFIHIPGERDLNLCIAPNPP
						AIFPTSIGYLGPDLNNLPTYAPFI
						TTNLTGPAPTTTHPVGNLGTTG
İ			i.			NPEPGGCGILIVWCCVNWACL
						WDSVVAAPREPYMPAGLLRHR
						RHYRHRREDLIAVPAARQTDE
						NRRPALCHEYAVGHRKIKTNV
		į				EHTEPYILPLITHSQPLNRGAFF
						CPGVRFLVVSVLDFEKRL**RR
						WRRSPAGI*GSRGAATTLSQRQ
						AQLTQHQTIRIPHPPGSGLPVVP
						KFPTGCVVVGAGPVRFVVING
24493	54861	Α	24631	1	924	
24494	54862	В	24632	177	468	
24495	54863	В	24633	592	883	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24496	54864	A	24634	1	1926	MATAKANQEIQEWLGKYGTAR
						VKLNVDKDFSLKDSSLEMLYPI
	ŀ	İ				YDTPTNMLFTQGAIHRTDDRTQ
						SNIGFGWRHFSGNDWMAAVIT
						FIDHDLSRSHTRIGVGAEYWRD
						YLKLSANGYIRASGWKKSPDIE
						DYQERPANGWDIRAEGYLPAW
						PQLGASLMYEQYYGDEVGLFG
						KDKRQKDPHAISAEHSPEFRSE
						ALKLVNASVLLPQPVNSACMN
ļ						HNSTTGAVNSKISRRLLNVNW
						RCLPRLHVSNASWQNGMKSWL
						SSKRPRHTSEAPEMKYVFIEKH
						QAEFSIKAMCRVLRVARSGWY
						TWCQRRTRINIAVVRRADGSRT
						SFVYTSYLAKVNEEWKNNVGT
						GSTVKWPIGLGGKGNDGIAAF
				Ì		VQRLPGAIGYVEYAYAKQNNL
						AYTKLISA*NEVCLY*KTSG*V
						QHQSNVPR/RSGWPAAAGIRGV
						SGGQG*TLL*YAAQMAPELPSS
				1		TPATWRK*TKSGKTTLV/HGST
				1		VKWPIGLGGKGNDGIAAFVQR
						LPGAIGYVEYAYAKQNNLAYT
						KLISADGKPVSPTEENFANAAK
						GADWSKTFAQDLTNQKGEDA
				·		WPITSTTFILIHKDQKKPEQGTE
						VLKFFDWAYKTGAKQANDLD
						YASLPDSVVEQVRAAWKTNIK
						DSSVNFDMANVFFRVTPGNFL
		1				VNREASHQRRVFAVRAGVDFI
						YRASSDGLRRCRDRRTRFKRIN
24497	54865	A	24635	1387	1639	ESKPLNLGRGSGFLLTRALSAQ
24491	34603	^	24033	1507	1039	PLWQSPDQHLTPQAGLRSRL*A
						QRQA*QKTASRVV*PKPLWAL
						VKATSEDISVVL*ESACQC
24498	54866	A	24636	1570	2516	IEVAMSIVVKNNIHWVGQRDW
24490	34800	^	24030	1370	2310	EVRDFHGTEYKTLRGSSYNSYF
						IREEKNVLIDTVDHKFSREFVQ
						NLRNEIDLADIDYIVINHAEEDH
						AGALTELMAQIPDTPIYCTANAI
	•					DSINGHHHHPEWNFNVVKTGD
						TLDIGNGKQLIFVETPMLHWPD
						SMMTYLTGDAVLFNYDAFGQH
						YCDEHLFNDEVDQTELFEQCQ
						RYYANILTPFSRLVTPKITEILGF
						NLPVDMIATSHGVVWRDNPTQ
						IVELYLKWAADYQEDRITIFYD
						TMVNNTPMMADAIAQGIA\ETG
						PRLAGENFNV\ARNDKHEDPDL
L	<u> </u>	<u> </u>	<u> </u>	<u></u>	<u></u>	LSFRSERGA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	(4)	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24499	54867	Α	24637	24	363	
24500	54868	A	24638	2	366	RATPRHIIVRFTKVEMKEKMLR AAREKGRVTLKGKPIRLTADLS AETLQARREW*QNLSAFACL*R ILFLLHL*SLVWLDMKCWVENS FL*EC*ILAPTLFWLVGFLPRDP LLV*WASL*G*PDLSLWLPLAF FPSFQLW*I*QLCVLELL
24501	54869	Α	24639	232	292	
24502	54870	Α	24640	4	911	
24503	54871	A	24641	2795	3226	SWWISFLMCCWIWFASILLRIFA SMFIKDIGLKFPFLVVSLPGFGI RMMLAS*NELGRIPSFSIGWNSF RRNGTSSSLYLW*NSAVNPSGP GLFLVGKLLVIASISEPVIGLFTD STSSWFSLGRVYVSRNLSISSRF SSLFA
24504	54872	Α	24642	1	3285	
24505	54873	Α	24643	1	4368	
24506	54874	A	24644	200	335	YALLNKDSSPWYPFSVPVLASN TRYLWPLSST**GSLSVWLSPD V
24507	54875	Α	24645	174	263	
24508	54876	Α	24646	1244	1354	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	i e	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	:			sequence		
24509	54877	Α	24647	194	2481	GMFRMNLMEPLTQCINAGHEM
						TKAIAIAQFNDDSPEARKITRR
						WRIGEAADLVGVSSQAIRDAEK
						AGRLPHPDMEIRGRVEQRVGY
ĺ						TIEQINHMRDVFGTRLRRAEDV
1					İ	FPPVIGVAAHKGGVYKTSVSVH
		İ				LAQDLALKGLRVLLVEGNDPQ
						GTASMYHGWVPDLHIHAEDTL
						LPFYLGEKDDVTYAIKPTCWPG
	}					LDIIPSCLALHRIETELMGKFDE
						GKLPTDPHLMLRLAIETVAHDY
						DVIVI\DSAPNLGIGTINVVCAA
						DVLIVPTPAELFDYTSALQFFD
						MLRDLLKNVDLKGFEPDVRILL
						TKYSNSNGSQSPWMEEQIRDA
						WGSMV\IKNVVRETNEVGKAA
						PMVDSLIARVGVMARGNAITLP
					i	VCGRDVKFTLEVLRGDSVEKTS
						RVWSGNERDQELLTEDALDDLI
						PSFLLTGQQTPAFGRRVSGVIEI
						ADGSRRRKAAALTESDYRVLV
						GELDDEQMAALSRLGNDYRPT
						SAYERGQRYASRLQNEFAGNIS
	•					ALADAENISRKIITRCINTAKLP
						KSVVALFSHPGELSARSGDALQ
						KAFTDKEELLKQQASNLHEQK
						KAGVIFEAEEVITLLTSVLKTSS
						ASRTSLSSRHQFAPGATVLYKG
						DKMVLNLDSPQSFSPGAGDHA
						NSPQSSSPSTGVHINSPQLSSLS
1						AGVHTISPQSSNPSVGVTALSHP
04510	64070		24640	1361	1604	DLVWVFTLTAHSRPVLVWVFT
24510	54878	A	24648	<u> </u>	1694 3852	MKLMETLNHCINAGHEMTKAV
24511	54879	Α	24649	1	3832	1
						AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMEIRGRVEQRVGYTIEQI
l l						NHMRDVFGTRLRRAEDVFPPVI
						GVAAHKGGVYKTSVSVHRAQ
						DLALKGLRVLLVEGNAPQGTA
						SVYHGWVPDLHIHAEAALLPF
ļ	,					YLGEKDDVTYAIKPTCWPGLDI
						IPSCLALPRIEPELMGK/FDEGRS
		1				FMPGPP/HHHPLGRSSQ*QVSNS
						LDRAAKGN*KPLCHC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24512	54880	A	24650	1	1720	MEKLLLENDSLSKDVSRLETVF
2 1312	3 1000	,	2 1030	•	1720	VPPEKHEKEIIALKSNIVELKKQ
					•	LSELKKKCGEDQEKIHALTSEN
						TNLKKMMSNQYVPVKTHEEV
}						KMTLNDTLAKTNRELLDVKKK
						FEDINQEFVKIKDKNEILKRNLE
		İ				NTQNQIKAEYISLAEHEAKMSS
					-	LSQSMRKVQDSNAEILANYRK
						GQEEIVTLHAEIKAQKKELDTIQ
					:	ECIKIMKRAPVIPKHTLNTQPVE
						DTSLSTPAAPMVDSLIARVGVM
						ARGNAITLPVCGRDVKFTLEVL
						RGDSVEKTSRVWSGNERDQEL
}						LTEDALDDLIPSFLLTGQQTPAF
						GRRVSGVIEIADGSRRRKAAAL
						TESDYRVLVGELDDEQMAALS RLGNDYRPTSAYERGQRYATR
						1
						PQNEFAGNISAMADAENISRKII
						TRCINTAKLPKSVVALFSHPGE
						LSARSVIDSAPNLGIGTINVVCA
ŀ						ADVLIVPTPAELFDYTSALQFFD
ļ						MLRDLLKNVDLKGFEPDVRILL
						TKYTNINASQSP\WMREKFGMA
1		1				WESMVLKNVVRETDEVGKGQI
	:					RMRADFV\QAIDQRSSTGAWRN
24512	Z 4001	<u> </u>	0.4651	1004	2271	ALSIWEPVCNEIFDRLIKPRSEIT
24513	54881	Α	24651	1094	2271	APFHRPHTKQSLQEQTDVVSIN
						EFLKQMAF*AQKLKE/WRTSAA
			;			SDITPAGFCSCRAIKDAEKPGRL
		1				PHPD\MEFRDRVDQ\RVVSPFGQ
						II\HMRNVFGPRLPRAKDVFPPV
ŀ						IGVAAHKGGVYKTSVSVHLAQ
		l				DLALKGLRVLLVEGN\DP\QGT
						ASMVSREGDPALHI\YQEAPFLP
						FYLGEKDDVTYAIKPTCCPGLN
						IIPSCLALPRIETELMGKFDEGK
						LPTDPHLMLRLAIETVAHAHDY
		Ì				DVIVIDSAPNLGIGTINVVCAAD
ł						V\LIVLTPVE\MFDYTSARQFFD
						MLRDLLKNA\DLKGFEPEVRIL
						LTKYSNSNGSQFP\WMEEQIRD
						AWGSMVLKNVVRETG\EVGKG
						\QIRMRT\VFEQAIDQRSSTDTSL
						STPAAPMVDSLIARVGVMASR
24514	54882	Α	24652	1	1415	

SEQ ID	1 -	Met	SEQ ID NO:	Nucleotide	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1 .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24515	54883	Α	24653	3008	8165	ILIPFFIGLQNRFTQFRLSETKEIT
						NPYAMRLYESLFQSSKTNGSGI
						VSLKIDWIIERYQLPQSYQRMP
						DFRRRFLQVCVNEINSRTPMRL
						SYIEKKKGRQTTHIAIRDAEKA
						GRLPHPDMEIRGRVEQRVGYTI
						EQINHMRDVFGTRLRRAEDVFP
						PVIGVAAHKGGVYKTSVSVHL
						AQDLALKGLRVLLVEGNDPQG
			:			TA\SMYHEWVQNLHIHAEDTLL
						PFYLGKKDDVTYAIKPTCWPGL
						DIIPSCLALHRI
24516	54884	Ā	24654	1	848	
24517	54885	Α	24655	2354	3640	NVICNQELVYSSRHQFAPGATV
						LYKGDKMVLNLDRSRVPTEYR
		ļ				YGVFLNVLLPATKYFQLHDPSL
						RNMELNYSLASEDLVLKGLRV
						LLVEGNDPQGTASMYHGWVP
						DLHIHAEDTLLPFYLGEKDDVT
						YAIKPTCWPGLDIIPSCLALHRI
						ETHLMGKFDEGKLPTDPHLML
						RLAIETVAHDYDVIVIDSAPNL
						GIGTINVVCAADVLIVPTPVE\L
						FDYASARQFFHMLRPLLKNVD
						LKGFEPDVRILLTKYSNSNGSQ
						SPWMEEQIRDAWGSMVLKNV
						KEFAGNISALADAENISRKIITR
						CINTAKLPKSVVVFFLTPVNYL
					-	PGDALQKAFTDKEELLKQQAS
Ì						NLHEQKKAGVIFEAEEVITLLTS
						VLKTSSASRTSLSSRHQFAPGA
						TVLYKGDKMVLNLDRSRVPTE
	- 100 6				0.50	CIEKIEAILKELEKPAP
24518	54886	A	24656	667	870	QIVSSEPKYTDFLSKIPFEQRVR
		ļ				DVQRNGDHTEGAVR*LSIAFFS
0.4510	54005	<u> </u>	0.4657	-	1006	ALLKLLAFFTEEVRAAAVVRFV
24519	54887	В	24657	1	1086	WINTER DIDNI IVINI CIVIZZII N
24520	54888	Α	24658	2	250	WRTKSLDIPPLIYPNGIVKNILR
						HLSQLLNVPITRPFSWILRPF*RP
						*RKISTSFIRPLFTFLPA**MLIRS
	<u></u>	<u> </u>		<u> </u>		SVWQ*KTVSW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24521	54889	A	24659	1	1608	MAETAVINHKKRKNSPRIVQSN
						DLTEAAYSLSRDQKRMLYLFV
						DQIRKSDGTLLYMTYRKPDGS
						GIVSLKIDWIIERYQLPQSYQRM
						PDFRRRFLQCINAGHEMTKAIAI
		İ				AQFNDDSPEARKITRRWRIGEA
						ADLLGVSSQAIRDAEKAGPLPH
						PDMENRGRVEQRGGYTIEPNN
						HMRDGVGTRWRRAEDVF\PPLI
						G\VAGHKGGR\YKTSVSVHLAQ
						DLALKGLRVLLVEGNDPQGTA
						SMYHGWVPDLHIHAEDTLLPF
						YLGEKDDVTYAIKPTCWPGLDI
						IPSCLALHRIETELMGKFDEGKL
						PTDPHLMLRLAIETVAHDYDVI
			,			VIDSAPNLGIGTINVVCAADVLI
						VPTPAELFDYTSALQFFDMLPD
		1				LVKNRDLKGVEPDMRKQRHR
						KAKDLLAQDPTAMTTDELLKE
						ATRCPICTAYLEKPMSLECECV
						i e
						FCLSCTNSLQKEPQGEGVLCPF
						CLEASQKNNIRLNRQLGRLVSH
						IKELEPKLKKILKMNFKDAEVP
						RFSSPKPCSSFFSTCYLSKNCSN
24522	54890		24660	1	1128	HPCYSPYPGSGWG
24523	54891	A	24661	1035	1208	SLIDRIEALGIFRFAAV*GSAAY
2-523	34071	1	24001	1033	1200	HGV*YERRRTFLCMLFGYFLSR
						RLESLLLPEVQS
24524	54892	A	24662	1	1316	MEIRGRVEQRVGYTIEQINHMR
24324	34072	^	24002	1	1310	DVFGTRLRRAEDVFPPVIGVAA
						HKGGVYKTSVSVHLAQDLALK
						GLRVLLVEGNDPQGTASMYHG
						WVPDLHIHAEDTLLPFYLGEKD
						DVTYAIKPTCWPGLDIIPSCLAL
						HRIETELMGKFDEGKLPTDPHL
						MLRLAIÉTVAHDYDVIVIDSAP
						NLGIGTINVALQNFQRYTGIQH VHRIGMAERMWCDRNRERHT
				•		
						VSSSGGNRLPNPGPDRSDLALK
						GLRVLLVEGNDPQGTASMYHG
						WVPDLHIHAEDTLLPFYLGEKD
			[DVTYAIKPTCWPGLDIIPSCLAL
		1				HRIETELMGKFDEGKLPTDPHL
						MLRLAIETVAHDYDVIVIDSAP
						NLVFRICFVIWAQERIDIKGFEP
						DVRILLTKYSNSNGSQSPWMEE
						QIRHAWGSMVLKN/VVRETDE
1	1	1	1			VGKDTGVRSKSIWCHRNSRWE

Sequence 09/540,217 codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence deletion, =possible nucleotide inservations codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptide sequence codon for peptid	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
GLHNADHVHKHLLGDNR IMQPSAHHPARQPEHDIAF SSSVSVHLAQDLALKGLR EGNDPQGTASMYHGWVP HAEDTLLPFYIDPHLMLRL VAHDYDVIVIDSAPNLGIG VCAADVLIVPTPAELFDYT QFFDMLRDLLKNVDLKGF VRILVVRVSQRMVVAGCII KLLNVLICFPFSVKNRFVIF DFECVIRF 24526 54894 A 24664 1 3216 MKLMETLNQCINAGHEMT AIAQFNDDSPEARKITRRW EAADLVGVSSQAIRDAEK. LPHPDMEIRGRVEQRVGY NHMRDVFGTRLRRAEDVF FGDFADFGTTIKQDFRLLG VDRLLQLSQGQAVKGNQI SLVKRKTTLAPNTQTASPR DSLMQLARQVSRL/VKRAI WPGLDIIPSC\LALPRIETEL FDEGKLPTDPHLMLRLAIE HDYDVIVIDSAPNLG 24527 54895 A 24665 1 419 MSSSDKDFDFSVANMVNF VFPMGKYYTQADKVLIPL HDGICEIHVAKYAEIFGLTT SKDIRQALKSFAGKEVVFY VGCRR*KRL*IFSLVYQTC/ QRALQCTYQPISHSLLYRV VYAVSA	NO:	1	hod	l .	codon for peptide		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
AIAQFNDDSPEARKITRRW EAADLVGVSSQAIRDAEK, LPHPDMEIRGRVEQRVGY' NHMRDVFGTRLRAEDVF FGDFADFGTTIKQDFRLLG VDRLLQLSQGQAVKGNQI SLVKRKTTLAPNTQTASPR DSLMQLARQVSRL/VKRAI WPGLDIIPSC\LALPRIETEL FDEGKLPTDPHLMLRLAIE HDYDVIVIDSAPNLG 24527 54895 A 24665 1 419 MSSSDKDFDFSVANMVNF VFPMGKYYTQADKVLIPL, HDGICEIHVAKYAEIFGLTS SKDIRQALKSFAGKEVVFY VGCRR*KRL*IFSLVYQTC, QRALQCTYQPISHSLLYRV VYAVSA	24525	54893	A	24663	504	1254	SYVKYFPHQPAQKYFQQIHSAI GLHNADHVHKHLLGDNRYPIW IMQPSAHHPARQPEHDIAFGKC SSSVSVHLAQDLALKGLRVLLV EGNDPQGTASMYHGWVPDLHI HAEDTLLPFY\DPHLMLRLAIET VAHDYDVIVIDSAPNLGIGTINV VCAADVLIVPTPAELFDYTSAL QFFDMLRDLLKNVDLKGFEPD VRILVVRVSQRMVVAGCIIAFID KLLNVLICFPFSVKNRFVIFHSV DFECVIRF
VFPMGKYYTQADKVLIPLA HDGICEIHVAKYAEIFGLTS SKDIRQALKSFAGKEVVFY VGCRR*KRL*IFSLVYQTCA QRALQCTYQPISHSLLYRV VYAVSA	24526	54894	A	24664		3216	MKLMETLNQCINAGHEMTKAI AIAQFNDDSPEARKITRRWRIG EAADLVGVSSQAIRDAEKAGR LPHPDMEIRGRVEQRVGYTIEQI NHMRDVFGTRLRRAEDVFPPG FGDFADFGTTIKQDFRLLGQTS VDRLLQLSQGQAVKGNQLLPV SLVKRKTTLAPNTQTASPRALA DSLMQLARQVSRL/VKRAPTC WPGLDIIPSC\LALPRIETELMGK FDEGKLPTDPHLMLRLAIETVA
24528 54896 A 24666 1 2598	24527	54895	A	24665	1	419	MSSSDKDFDFSVANMVNFFAP VFPMGKYYTQADKVLIPLAIQE HDGICEIHVAKYAEIFGLTSAEA SKDIRQALKSFAGKEVVFYRPE/ VGCRR*KRL*IFSLVYQTCAQSI QRALQCTYQPISHSLLYRVTEP VYAVSA
124320 34030 1x 24000 1 2370	24528	54896	A	24666	1	2598	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24529	54897	Α	24667	1	2027	MRGHREKAAACKPRTRASEGT
,						TPASTFILDFQPPNCNYTSGDNH
						TLRDPHYVEDKGHKYLVFEAN
		1				TGTENGYQGEESLFNKAYYGG
		1				GTNFFRKESQKLQQSAKKRDA
						ELANGALGIIELNNDYTLKKVM
						KPLITSNTVTDEIERANVFKMN
						GKWYLFTDSRGSKMTIDGINSN
		l				DIYMLGYVSNSLTGPYKPLNKT
						GLVLQMGLDPNDVTFTYSHFA
						VPQAKGILCEEDNYTAGDNHM
						MRAPHCEEDRAHKFVVFDANA
						GTESGHQGESSLFNRACGGGGT
						FFFSKESQKLQQSAKKRDCINA
İ						GHEMTKAIAIAQFNDDSPEARK
						ITRRWRIGEAADLVGVSSQAIR
						DAEKAGRLPHPDMEIRGRVEQ
						RVGYTIEQINHMRDVFGTRLKR
						AEDVFPPVIGVAAHKGGVYKT
	ļ					SVSVHLAQDLALKGSLLLPKND
						FLFKLGLEGHLPLGKIHSPPTGA
						DVARGSSGLPKSELFLPERNTQ
			ļ			ELQQDSEEGPLALQVLQSNLM
						DFADFGTTIKQDFRLLGQTSVD
		ĺ				RLLQLSQGQAVKGNQLLPVSL
İ		ļ				VKRKTTLAPNTQTASPRALADS
						LMQLARQVSRLESGQNNDGIC
						EIHVAKYVEIFGLTSAEASKDIR
ŀ						QALKSFAGKLVVFYRPE/VGCR
						R*KRL*IFSLVYQTCAQSIQRAL
						QCTYQPISHSLLYRVTEPVYAV
24530	54898	Α	24668	1	989	PWISAPVPVDVVEGAMDSVTV
		1				LSFGGLMLYFCAGWPPARRWC
						FPESISCGSMERDQWWGLQVA
						KRAGLAGGQSGRTVLRERVRIE
		}				IASTHIALAARHSDWRCCRNGR
						YPARGPAALQNFQRYTGIQHV
						HRIGMAERMWCDRNRERHTVS
		1				SSGGNRLPNPGPDRSCDNLKTC
		1				HTSHGSVMAETAVINHKKRKN
		1				SPRIVQSNDLTEAAYSLSRDQK
]			RMLYLFVDQIRKSDGTLQEHD
						GICEIHVAKYAEIFGLTSAEASK
						DIRQALKSFAGKEVVFYRPE/V
						GCRR*KRL*IFSLVYQTCAQSIQ
						RALQCTYQPISHSLLYRVTEPV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24531	54899	Α	24669	350	1203	LLTARLTWVSARLMSDFADFG
						TTIKQDFRLLGQTSVDRLLQLS
						QGQAVKGNQLLPVSLVKRKTT
						LAPNTQTASPRALADSLMQLA
						RQD/LALKGLRVLLVEGNDPQG
						TASMYHGWVPDLHIHAEDTLL
						PFYLGEKDDVTYAIKPTCWPGL
1						DIIPSCLALHRIETELMGKFDEG
						KLPTDPHLMLRLAIETVAHDYD
						VIVIDSAPNLGIGTINVVCAADV
		1				LIVPTPAELFDYTSALQFFDMLR
						DLLKNVDLKGFEPDVRILLTKY
		Ì				
2.4522	54000	ļ	0.4670	1	026	SNSNGSQSPWMEEQIRDAEGSR
24532	54900	A	24670	1	936	L CTTPG C COPN TTG LL L VOI VOI VOI TO
24533	54901	Α	24671	1070	1324	LGTPGGCSRMTCALVPLIPNDD
		l				TPTRRTRPLLGQGVFSVRS*TVP
						ADQSTSGDGDSTCNVRGSCPCR
						IAITILITPATPAAACV
24534	54902	Α	24672	1	1212	
24535	54903	Α	24673	2373	3381	FKRGQYGQLLRPRFHHGQILYA
						RRQGADAAGDSGIW/HGIWTEP
		1				MTM*KKDER*PSPTQAS*DNW
		ļ				KSIRGSCWQALSITRN/CKNSPRI
		İ				VQSNDLTEAAYSLSRDQKRML
						YLFVDQIRKSDGTLQEHDGICEI
		İ			1	HVAKYAEIFGLTSAEASKDIRQ
						ALKSFAGKEVVFYRPEEDAGD
						EKGYESFPWFIKRAHSPSRGLY
						SVHINPYLIPFFIGLQNRFTQFRL
						SETKEITNPYAMRLYESLCQYR
						KPDGSGIVSLKIDWIIERYQLPQ
		1				SYQRMPDFRRRFLQHIFVLRER
						PETVLIDLIQRTKDAVRELDNL
		1				QYRKMKKLLFQEAHNGPAVEA
		1				QEEEEDD
24536	54904	Α	24674	31	811	RRGLLGTRLRLSEDVFPPVLGV
						AAHKGGVDKTSVSVHLAQDLA
		1				LKGLRVLLVEGNDPQGTASMY
						HGWVPELHIHAEDTLLPFYLGE
						KDDVTYAIKPTCWPGLDIIPSCL
		1				ALHRIETELMGKFDEGKLPTDP
		1				HLMLRLAIETVAHDYDVIVIDS
						APNLGIGTINVVCAADVLIVPTP
		İ				AELFDYTSALQFFDMLRDLLKN
						VDLKGFEPDVRILLTKYSNSNG
	1	1				1
		[SQSPWMEEQIRDAWG\SMVLK
0.4.50=	54005	 	04675	1004	1105	NVVAFQNLEFLFILLRIWL
24537	54905	A	24675	1004_	1105	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24538	54906	A	24676		1139	MEKKITGYTTVDISQWHRKEHF EAFQSVAQCTYNQTVQLDIKK KLKTVKKNKHKFYPAFIHILAR LMNAHPEFRMAMKDVAEYNG YHVVFALAGSPKDADDTSIYM FYQKVGDNSIDSWKNAGRVFK DSDKFDANDPILKDQTQEWSGS ATFTSDGKIRLFYTDYSGKHYG KQSLTTAQVNVSKSDDTLKING VEDHKTIFDGDGKTYQNVQQFI DEGNYTSGDNHTLRDPHYVED KGHKYLVFEANTGTENGYQGE ESLFNKAYYGGGTASSIRKSDG TLQEHDGICEIHVAKYAEIFGLT SAEASKDIRQALKSFAGKEVVF YRPE/VGCRR*KRL*IFSLVYQT CAQSIQRALQCTYQPISHSLLYR
						VTEPVYAVSA
24539	54907	A	24677	1	930	
24540	54908	A	24678		2486	MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQNEKY QVPQFDQSTIKNIESAKGLDVW DKWPLANADGTVAEYNGYNV VDKFDANDPILKDQTQEWSGS ATFTCDGKIRLFYTDYSGKHYG KQSLTTAQVNVLKSDDTLKING VEDHKAIFDGDGKTYQNVQQL IDEGNYTSGDNHTLRDPHYVED KGHKYLVFEANTGTENGYQGE ESLFNKAYYGGGTNFFRKESQK LQQSAKKRDAELANGALGIIEL NNDYTLKKVMKPLITSNTVTDE IERANVFKMNGKWYLFTDSRG SKMTIDGINSNDIYMLGYVSNS LTGPYKPLNKTGLVLQMGLDP NDVTFTYSHFAVPQAKGNNVV ITSYMTNRGFFEDKKATFAPSF LMNIKGNKTSVVKNSILEQGQL TWLQVAKRAGLGGGQSGRTVL RERLPNYKNFKGTIQELGQNQY AVSGEIFVVDRNTVEITELPVRT WTQVYKEQVLEPMLNGTDKTP ALISDYKEYHTDTTVKFVVKM TEEKLAQAEAAGLHKVFKLQT TLTCNSMVLFDHMGCLKKYET VQDILKEFFDLRLSYYGLRKEW LVGMLGAESTKLNNQARFILEK IQGKITIENRSKKDLIQMLVQRG YESDPVKAWKEAQEKCDNLKT CHTSHGSVMAETAVINHKKRK

SEQ ID	SEQ ID NO:		1		j .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24541	54909	A	24679	5	253	SAPKVWRGLRQQGAGISVLP*V CAQP/VQVAKAPELILNFDLPK KYFDKL/R/DTHFFSPHTGKFPT VKILMAPFANSASRGVALC
24542	54910	Α	24680	334	842	PDKEGNEIWVDMYTVKPSGWT VRTFDKPRKRFIAFFIAGILFRAI KNHFLPRETLQCLPYILTGFRRG QSEYFSIFSNMDLADTVMFLVG GHKIFVFHCHTELRMSIHQAGK NVNKGRIKLVLIFLYGL*/QRP* YPAERSGYRYIEQLTEMPQNVL YDAIGIYQRWYIQ
24543	54911	A	24681	676	1169	DNLKTCHTSHGSVMAETAVIN HKKRKNSPRIVQSNDLTEAAYS LSRDQKRMLYLFVDQIRKSDGT LQEHDGICEIHVAKYAEIFGLTS AEASKDIRQALKSFAGKEVVFY RPE/VGCRR*KRL*IFSLVYQTC AQSIQRALQCTYQPISHSLLYRV TEPVYAVSA
24544	54912	Α	24682	1	783	
24545	54913	Α	24683	1	1884	
24546	54914	Α	24684	1	1131	
24547	54915	Α	24685	1	2856	
24548	54916	Α	24686	528	725	
24549	54917	A	24687	1	898	
24550	54918	A	24688	1879	2343	
24551	54919	A	24689	1	1254	
24552	54920	A	24690	1	1113	
24553	54921	В	24691	1	1156	
24554	54922	В	24692	1	1980	
	54923	В	24693	1	1932	
24555 24556	54924	A	24694	1	663	
24557	54925	A	24695	1123	1764	
24558	54926		24696	2	1772	
24559	54927	A	24697	1	1040	MELKTKARELREECRSLRSRCN
						QLEERVSAMEDEMNEMKREG KFREKRIKRNEQSLQEIWDYVK RPNLRLIGVPESDVENGTKLEN TLQDIIQENFPNLARQANVQIQE IQRTPQRYSSGRETPRHIIVRFT KVEMKEKMLRAAERKCDNLK TCHTSHGSVMAETAVINHKKR KNSPRIVQSNDLTEAAYSLSRD QKRMLYLFVDQIRKSDGTLQE HDGICEIHVAKYAEIFGLTSAEA SKDIRQALKSFAGKEVVFYRPE EDAGDEKSYEQPSKDIRQALKS FAGKEVVFYRPE/VGCRR*KRL *IFSLVYQTCAQSIQRALQCTYQ PISHSLLYRVTEPVYAVSA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
i	sequence		09/340,217	sequence	or peptide sequence	deterior, —possible nucleonide insertion)
24560	54928	IA	24698	1683	4289	 RVFKDSDKFDANDPILKDQTQE
2.000	1000	` `		1.000		WSGSATFTSDGKIRLFYTDYSG
		İ				KHYGKQSLTTAQVNVSKSDDT
		İ				LKINGVEDHKTIFDGDGKTYQN
		1		1		VQQFIDEGNYTSGDNHTLRDPH
						YVEDKGHKYLVFEANTGTENG
						YQGEESLFNKAYYGGGTNFFR
						KESQKLQQSAKKRDAELANGA
						LGIIELNNDYTLKKVMKPLITSN
						TVTDEIERANVFKMNGKWHSK
						ILVSGTHRARTEVEVTDPIHQQ
						RMGCCFTVGAGEICEKPTGCFF
						TPLPNCNCQWTGSAILDRRSHS
						HPHSPASGSWVKVVTGSYQLT
						HSPGIGLRDFADFGTPHQTGFSP
		1		1		AGANQRGPLVATLSGPGGEGQ
						SAVARLTGEKKNHPGAQYANR
						LSPRVGRFINAAGTTGFPTGKR
		1				AVSATQLMHFKILVSRAHRART
	1					ELEVTDPGSSAENGLLFPVVLL
						LRKARKTLMPPHSTCFIKRSAP
						TQSTAGKTRPRVLKDSDKFDA
						NDPILKDQTQEWSGSAPFTSDG
				,		KIRLFYTDYSGKHYGKQSLTTA
		1				QVNVSKSDDTLKINGVEDHKTI
		1				FDGDGKTYQNVQQFIDEGNYT
						SGDNHTLRDPHYVEDKGHKYL
1						VFEANTGTENGYQGEESLFNKS
						VLRRRHELLPDFADFGPTIKQD
						FRLLGQTSVDRLLQLSQGQAV
			-			KGNQLLPVSLVKRKTTLAPNTQ
24561	54929	В	24699	1	2034	TASPRALADSLMQLARQVSRLE
24562	54930	A	24700	1	1038	
24563	54931	A	24701	1	1149	
24564	54932	A	24702	2	823	WPLQNADGTVAEYNGYHVVF
		1				ALAGSPKDADDTSIYMFYQKV
		1				GDNSIDSWKNAGRVFKDSDKF
						DANDPILKDQTQEWSGSATFTS
						DGKIRLFYTDYSGKHYGKQSLT
						TAQCDNLKTCHTSHGSVMAET
						AVINHKKRKNSPRIVQSNDLTE
						AAYSLSRDQKRMLYLFVDQIR
						KSDGTLQEHDGICEIHVAKYAE
						IFGLTSAEASKDIRQALKSFAGK
						EVVFYRPEEDAGDEK/ML*IFSL
!						VYQTCAQSIQRALQCTYQPISH
	<u></u>			<u></u>		SLLYRVTEPGYAVSA
24565	54933	В	24703	1	1206	
24566	54934	Α	24704	2	1037	
24567	54935	В	24705	1	1389	

SEQ ID			SEQ ID NO:	Nucleotide	ID .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	or peptide sequence	deletion, (=possible nucleotide insertion)
24568	54936	Α	24706	1	602	MGTGDFICISMTGGAPWGFRLQ
						GGKEQKQPLQVAKWANLKPGP
		1		:		PSHGSVMAETAVINHKKRKNSP
						RIVQSNDLTEAAYSLSRDQKRM
						LYLFVDQIRKSDGTLQEHDGIC
						EIHVAKYAEIFGLTSAEASKDIR
					1	QALKSFAGKEVVFYRPE/VGCR
						R*KRL*IFSLVYQTCAQSIQRAL
		:				QCTYQPISHSLLYRVTEPVYAV
24569	54937	Α	24707	1	774	
24570	54938	A	24708	1479	1947	
24571	54939	Α	24709	1	696	
24572	54940	A	24710	1	1773	
24573	54941	A	24711	1	1509	
24574	54942	A	24712 24713	156	891 242	VAGRGIGYRLSPNKGL*TWSAL
24575	54943	Α				CKPIAE CKPIAE
24576	54944	Α	24714	1441	1866	
24577	54945	Α	24715	1	1098	
24578	54946	A	24716	1	1989	
24579	54947	В	24717	126	848	
24580	54948	В	24718	58	688 542	
24581 24582	54949 54950	A A	24719 24720	1	675	
24583	54951	A	24721	1	1209	MKMPEKHDLLAAILAAKEQGI
24363	34931	^	24/21	1	1207	GAILAFAMAYLRGRYNGVLEA
		l				RSLKSRCQQSHALAEGSRGGPP
			,			VTGRFKSTFTADSWNRYSLPSL
			}			TGGGNILHSRTHRRTVSRLLGR
	1					RWQELLRDVFGTRLRRAEDVF
						PPVIGVAAHKGGVYKTSVSVH
						LAODLALKGLRVLLVEGNDPO
						GTASMYHGWVPDLHIHAEDTL
						LPFYLGEKDDVTYAIKPTCWPA
						LDIIPSCLALHRIETELMGKFDE
						GKLPTDPHLMLRLAIETVAHDY
						DVIVIDSAPNLGIGTINVVCAAD
						VLIVPTPAELFDYTSALQFFDM
						LRDLLKNVDLKGFEPDVRILLT/
						KYSNSNGSQSPWMEEQ\IRDAW
						GSMGLKNV/VRETDEIG\KGQIR
				1		MRTVFETAMDKRLLTGAWEK
						AFFFLEPGFEEKSNR

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24584	54952	A	24722		743	MIGMELPGLEGTELEGCHEREV VYDRVRVFEKRGEARHVEARS PGGGSEKRGWGGARHQRMCR KGADNGGAGRARRVARLRQL ASRAIGRRKGAVLYQTPVINHK KRKNSPRIVQSNDLTEAAYSLS RDQKRMLYLFVDQIRKSDGTL QEHDGICEIHVAKYAEIFGLTL WREASKDIRQALKSFAGKEVV FYRPE/VGCRR*KRL*IFSLVYQ TCAQSIQRALQCTYQPISHSLLY RVTEPVYAVSA
24585	54953	Α	24723	1157	1471	
24586	54954	A	24724		824	MEENERDLEINMINMSKELKVE NIKKHIGMLEKDEIEIIVTKSILI EIEIPTQCLRNLNTLHLAASAGN HMSNTACWYGCEDFTSVSTFA HQGVGSWDAYEPAASLLLSCD NLKTCHTSHGSVMAETAVINH KKRKNSPRIVQSNDLTEAAYSL SRDQKRMLYLFVDQIRKSDGTL QEHDGICEIHVANYAEIYGLTS AEASKDIRQALKSFAGKEVVFY RPE/VGCRR*KRL*IFSLVYQTC AQSIQRALQCTYQPISHSLLYRV TEPVYAVSA
24587	54955	A	24725	996	1169	SLIDRIEALGIFRFAAV*GSAAY HGV*YERRRTFLCMLFGYFLSR RLESLLLPEVQS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24588	54956	Α	24726	3	1743	DSPE\ARKITRRWRIGEAADLV
	ļ	ŀ				GVSSQAIRDAEKAGRLPHPDME
		ľ				IRGRVEQRVGYTIEQINHMRDV
1		ļ				FGTRLRRAEDVFPPVIGVAAHK
	i					GNDPQGTASMYHGWVPDLHIH
1						AEDTLLPFYLGEKDDVTYAIKP
		İ				TCWPGLDIIPSCLALHRIETELM
		ŀ				GKFDEAQPNLGIGTINVVCAAD
		ŀ				VLIVPTPAELFDYTSALQFFDM
						LRDLLKNVDLKGFEPDVRILLT
						KYSNSNGSQSPWMEEQIRDAW
İ						GSMVLKNVVRETDEVGKGQIR
						MRTVFEQAIDQRSSTDTSLSTP
						AAPMVDSLIARVGVMARGNAI
						TLPVCGRDVKFTLEVLRGDSVE
						KTSRVWSGNERDQELLTEDAL
		ŀ				DDLIPSFLLTGQQTPAFGRRVSG
						VIEIADGSRRRKAAALTESDYR
		ļ				VLVGELDDEQMAALSRLGNDY
	<u> </u>	ŀ				RPTSAYERGQRYASRLQNEFAG
						KYFLRWLMRKIFHIITRCINTAK
		ļ				LPKSVVALFSHPGELSARSGDA
						LQKAFTDKEELLKQQASNLHE
		ŀ				QKKAGVIFEAEEVITLLTSVLKT
						SSASRTSLSSRHQFAPGATVLY
		•				KGDKMVLNLDRSRVPTECIEKI
		İ				EAILKELEKPAP
24589	54957	A	24727	1	1623	
24590	54958	Α	24728	1	1863	
24591	54959	Α	24729	1	2622	
24592	54960	Α	24730	1	1338	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Į.	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24593	54961	A	24731		1823	MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGVED HKTIFDGDGKTYQNVQQFIDEG NYTSGDNHTLRDPHYVEDKGH KYLVFEANTGTENGYQGEESLF NKAYYGGGTNFFRKESQKLQQ SAKKRDAELANGALGIIELNND YTLKKVMKPLITSNTVFIRDKL MERRNRTGRTEKARIWEVTD RTVRTWIGEAVAAAAADGVTF SVPVTPHTFRHSYAMHMLYAG IPLKVLQSLMGHKSISSTEVYTK VFALDVAARHRCDNLKTCHTS HGSVMAETAVINHKKRKNSPRI VQSNDLTEAAYSLSRDQKRML YLFVDQIRKSDGTLQEHDGICEI HVAKYAEIFGLTSAEASKDIRQ ALKSFAGKEVVFYRPE/VGCRR
24594	54962	A	24732	1476	2221	*KRL*IFSLVYQTCAQSIQRALQ CTYQPISHSLLYRVTEPVYAVS

SEQ ID		Met	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24595	54963	Α	24733	1	815	MRVGRKAVGAIWWSSKQELFQ
24393	34903		24733	1	013	QLFLRTELEDIGPGIVKRELGRE
						GSKDTGSSDDWEQLPDWKTKR
						1
						VGNERDQELLIEDALDDLIPSFL
						LTGQQTPAFGRRVSGVIEIADG
						SRRRKAAALTESDYRVLVGEL
		ŀ				DDEQMAALSRLGNDYRPTSAY
						ERGQRYASRLQNEFAGNISALA
						DAENISRKIITRCINTAKLPKSV
						VALFSHPGELSARSGDALQKAF
		ŀ				TDKEELLKQQASNLHEQKKAG
	ŀ					VIFEAEEVITLLTSVLKTVQSKN
						LSHFTWICHAETAVINHKKRKN
						SPRIVQSNDLTEAAYSLSRDQK
						RMLYLFVDQIRKSDGTLQEHD
						GICEIHVAKYAEIFGLTSAEASK
						DIRQALKSFAGKEVVFYRPEED
						AGDEKGYESFPWFIKHWVTISP
					[QTSAYERGQRYASRCKNEFAG
					1	NISALADAENISRKIITRCINTAK
	1					LPKSVVALFSHPGELSARSGDA
						LQKAFTDKEELLKQQASNLHE
						QKKAGVIFEAEEVITLLTSVLKT
						SSASRTSLSSRHQFAPGATVLY
						KGDKMVLNLDRSRVPTECKRK
						LRPF*SGQRTDTAYVFDPGRDY

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24596	54964	A	24734	2	1249	DFADFGTTIKQDFRLLGQTSVD RLLQLSQGQAVKGNQLLPVSL VKRKTTLAPNTQTASPRALADS LMQLARQVSRLESGQLAISGRI LFLPMLNSINKLAAPSQPCTEPF FRRDPTLHPKSEVRLPPKVTTIT VISNPILITPQEKSVAQCTYNQT VQLDITAFLKTVKKNKHKFYPA FIHILARLMNAHPEFRMAMKD GELVIWDSVHPCYTVFHEQTET FSSLWSEYHDDFRQFLHIYSQD VACYGENLAYFPKGFIENMFFV SANPWVSFTSFDLNVANMDNF FAPVFTMGKYYTQGDKVLMPL AIQIRKSDGTLQEHDGICEIHVA KYAEIFGLTSAEASKDIRQALKS FAGKEVVFYRPEEDAGDEKGY ESFPWFIKRAHSPSRGLYSVHIN PYLIPFFIGLQNRFTQFRLSETKE ITNPYAMRLYESLCQYRKPDGS GIVSLPIDWIIERYQLPQSYQRM PDFRRRFLQVCVNEINSRTPMR LSYIEKKKGRQTTHIIRISDGTL QEHDGICEIHVAKYAEIFGLTSA EASKDIRQALKSFAGKEVVFYR PEEDAGDERGYESFPWFIKRAH SPSRGLYSVHINPYLIPFFIGLQN RFTQFRLSETKEITNPYAMRLY ESLCQYHLCGTLRNSRFCEIHV AKYAEIFGLTSAEASKDIRQAL
						KSFAGKEVVFYRPE/VGCRR*K RL*IFSLVYQTCAQSIQRALQCT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24597	54965	A	24735	I	2286	MLTLILLDLQFMAQLAGYQMS FNSTKSKQTKNNFAVGYRTGD FQLHTNVNDGAEFGGSVYQKV CEELDALINLAWTSSTSCTRFG LAAKFQLKPIAFISTKVNHWLT GVSYTPPLEAGVKLNLSALCDN LKTCHTSHGSVMAETAVINHK KRKNSPRIVQSNDLTEAAYSLS RDQKRMLYLFVDQIRKSDGTL QEHDGICEIHVAKYAEIFGLTSA EASKDIRQALKSFAGKEVVFYR PEEDAGDEKGYESFPWFIKRAH SPSRGLYSVHINPYLIPFFIGLQN RFTQFRLSETKEITNPYAMRLY EIPVVSIRKPGMGSGIVSLKIDW IIERYQLPQSYQRMPDFRRFLQ CINAGHEMTKAIAIAQFNDDSP EARKITRRWRIGEAADLLGVSS QAIRDAEKAGPLPHPDMENRG RVEQRGGYTIEPNNHMRDGVG TRWRRAEDVF\PPLIG\VAGHKG GR\YKTSVSVHLAQDLALKGLR VLLVEGNDPQGTASMYHGWV PDLHIHAEDTLLPFYLGEKDDV
						TYAIKPTCWPGLDIIPSCLALHR IETELMGKFDEGKLPTDPHLML RLAIETVAHDYDVIVIDSAPNL GIGTINVVCAADVLIVPTPAELF DYTSALQFFDMLPDLVKNRDL KGVEPDVFIRDKLMERRNRRT GRTEKARIWEVTDRTVRTWIGE AVAAAAADGVTFSVPVTPHTF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24598	54966	A	24736	11	1400	MDAKMHSGGTAPPGTLSQRQP
			İ			SSNSKTNGSKLNPLKDTLLSTPS
						CPEGELVNAARRSNGSGNAITL
						PVWGRDVKFTLEVLRGDSVEK
		İ				TSRVWSGNERDQELLTEDALD
				}		DLIPSFLLTGQQTPAFGRRVSGV
						IEIADGSRRRKAAALTESDYRV
		ŧ				LVGELDDEQMAALSRLGNDYR
		ļ				PTSAYERGQRYASRLQNEFAG
						NISALADAENISRKIITRCINTAK
				·		LPKSVVALFSHPGELSARSGDA
						LQKAFTDKEELLKQQASNLHE
						QKKAGCDNLKTCHTSHGSVMA
						ETAVINHKKRKNSPRIVQSNDL
						TEAAYSLSRDQKRMLYLFVDQI
						RKSDGTLQEHDGICEIHVAKYA
[İ		EIFGLTSAEASKDIRQALKSFAG
İ						KEVVFYRPEEDAGDEKSYEQPS
						KDIRQALKSFAGKEVVFYRPE/
						VGCRR*KRL*IFSLVYQTCAQSI
ŀ						QRALQCTYQPISHSLLYRVTEP
24599	54967	A	24737	1	1628	MLHITNDQGNASQNHNAMPPY
						SCKNGHNLKTVDGGMDAVNG
						EHFYTAGLSQGPLKGSFLLFFNI
					ļ	LILVCPPIFRVSGNEVFIRDKLM
		İ				ERRNRRTGRTEKARIWEVTDRT
						VRTWIGEAVAPLLLTCDNLKTC
						HTSHGSVMAETAVINHKKRKF
ļ						SPRIVQSNDLTEAAYSLSRDQK
		1				RMLYLFVDQIRKSDGTLQEHD
						GICEIHVAKYAEIFGLTSAEASK
	:	1				DIRQALKSFAGKEVVFYRPEED
						AGDEKGYESFPWFIKRAHSPSR
		İ				GLYSVHINPYLIPFFIGLQNRFT
						QFRLSETKEITNPYAMRLYESL
						CQYRKPDGSGIVSLKIDWIIERY
						QLPQSYQRMPDFRRRFLQVCV
	1					NEINSRTPMRLSYIEKKKGRQT
						THICDNLKTCHTSHGSVMAETA
						VINHKKRKNSPRIVQSNDLTEA
						AYSLSRDQKRMLYLFVDQIRKS
						DGTLQEHDGICEIHVAKYAEIF
						GLTSAEASKDIRQALKSFAGKE
		1				VVFYRPE/VGCRR*KRL*IFSLV
						YQTCAQSIQRALQCTYQPISHSL
						LYRVTEPVYTISAY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24600	54968	Α	24738	1295	1675	SVTSCLKVTSVCLTSGRRFLQIR
						ISDGTLQEHDGICEIHVAKYAEI
		1				FGLTSAEASKDIRQALKSFAGK
				ŀ		EVVFYRPEEDAGDEKGYESFP
						WFIKRAHSPSRGLYSVHINPYLI
						PFFIGLQNRFTQFRLSETKEITNP
					ļ	YAMRLYESLCQYRKPDGSGIVS
	ļ			ļ		LKIAWVIERYQLPQSYQRMPDF
		l				RRRFLQVSDFLMAPYRNIDGIC
		1				EIHVAKYAEIFGLTSAEASKDIR
		1				QALKSFAGKEVVFYRPEEDAG
						DEKGYESFPWFIKRAHSPSRGL
		ŀ				YSVHINPYLIPFFIGLQNRFTQFR
	İ					LSETKEITNPYAMRLYESLCQY
						RKPDGSGIVSLKIDWIIERYQLP
						QSYQRMPDFRRRFLQVCVNEIN
		ŀ				SRTPMRLSYIEKKKGRQTTHIE
ļ						HDGICEIHVAKYAEIFGLTSAEA
						SKDIRQALKSFAGKEVVFYRPE
				:		EDAGDERGYESFPWFIKRAHSP
						SRGLYSVHINPYLIPFFIGLQNRF
						TQFRLSETKEITNPYAMRLYQS
						LCQYRKPDGSGIVSLKIDWIIER
ŀ						YQLPQSYQRMPDFRRRFLQIRK
						SDGTLQEHDGICEIHVAKYAEIF
						GLTSAEASKDIRQALKSFAGKE
						VVFYRPEEDAGDEEGYESFPWF
						IKRAHSPSRGLYSVHINPYLIPFF
						IGLQNRFTQFRLSETKEITNPYA
1					}	MRLYESLCQYRKPDGSCIVSLK
1						IDWIIERYQLPPRYQRMPDFRRR
						FLRVWVQEINSRTPMPPLLPWE
24601	54969	Α	24739	1	1986	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon,/=possible nucleotide
	sequence	,	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24602	54970	A	24740	1	398	MVDSLIARVGVMARGNAITLP
						VCGRDVKFTLQVLRGDSVEKT
						SRVPPGNERDQEMLTEDALDD
			-			LIPSFLLTGQQTPAFGRRVSGVI
			ļ			HTADGSRRRKAAALTEKFCIVD
						NPPGEIRKRACLFIPTHALNLNR
						FEDICYAHQLPDGGLLSARVGV
						MARGNPILSPVCGRNVKFTLKV
						LRGDSVENPSRVWSGNERAQE
						LLTEHALDDLTLSFLLTGQQTP
						AFGRRVSGVIEIADGSRRRKAA
						ALTESDYRVLVGELDDEQMAA
						LSRLGNDYRPTSPYERGQRYAS
						RLQNEFAGNISALADAENISRKI
						ITRCINTAKLPKSVVALFSHPGE
		l				LSARSGNERDQELLTEDALDDL
						IPSFLLTGQQTPAFGRRVSGVIH
						TADGSRRRKAA/DTYRKILYR**
						PPGGD*KKSVPFYSNTCAQSQP
24603	54971	Α	24741	1450	3699	RNPLQIPKQQQNEKYQVPQFDQ
						STIKNIESAKGLDVWDSWPLQN
						ADGTVAEYNGYHVVFALAGSP
						KDADDTSIYMFYQKVGDNSIDS
						WKNAGRVFKDSDKFDANDPIL
						KDQTQEWSGSATFTSDGKIRLF
		1				YTDYSGKHYGKQSLTTAQIFQ
						MITVFITSLCNTTSSHICLHISWV
				į		TTMHGAMPHPGHIPTAPASLPA
						KVKGSGKVCWLAFPLSELTPPS
						ALQVNVSKSDDTLKINGVEDH
						KTIFDGDGKTYQNVQHFIDEGN
						YTCGDNHTLRDPHYVEDKGHK
		1				YLVFEANTGTENGYQGEESLFN
		1				KAYYGGGTNFFRKESQKLQQS
						AKKRDAELANGALGIIELNNDY
		l				TLKKVMKPLITSNTVTDEIERA
						NVFKMNGKWYLFTDSRGSKM
		1				TIDVPQAKSTNVVITSYMTNRG
		İ				FFEPEGSKGSITAAVMAKGLSV
						ASGNSTPERRRATANENVQPRA
						IRDAEKAGRLPHPDMEIRGRVE
						QRVGYTIEQINHMRDVFGTRLR
		1				RAEDVFPPVIGVAAHKGGVYK
						TSVSVHLAQDLALKGLRVLLV
						EGNDPQGTASMYHGWVPDLHI
		1				HAEDTLLPFYLGEKDDVTYAIK
						PTCWPGLDIIPSCLALHRIETEL
		1				MRKFDEGNLPTDPHLMLRLAIE
						TVAHDYDVIVIDSAPNLGIGTIN
		1				VVCAADVLIVPTPAEQLDIS\SS
L		<u>l</u>		<u> </u>		RQFFD\ILRDLLKNVDLKGFEPD

-	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24604	54972	Α	24742	1	3899	MVDSLIAPVGVMARGNAITLPV
						RATNVKFTLEELRGDSVEKTSR
						VWSEKKDRGGQEAEVSGEGSS
						QTKSLQKRQHSNGSDFTARGF
						HTGQALVRSKGTYWNPTGLGL
						SHGTEDPNLIGEELDAGEWDK
					·	NTRKELAVLNVRRDFADFGTTI
						KQDFRLLGQTSVDRLLQLSQG
						QAVKGNQLLPVSLVKRKTTLA
						PNTQTASPRALADSLMQLARQ
						VSRLESGQFWVFEGGLRVLVR
						ERVTTPKGT\ASMYQGW\VPD
24605	54973	Α	24743	268	2617	RMFRMKLMETL\SQCINAGHE
						MTKAIAIAQFNDDSPEARKITR
						RWRIGEAADLVGVSSQAIRDAE
						KAGRLPHPDMEISGR\VKHGVG
						YTIEQIYL\TRDVFGTRLPRAED\
						VFPPVIGVAAHKGGVYKTSVSV
						HLAQDLALKGLRVLLVEGNDP
						QGTASMYHGWVPDLHIHAEDT
						LLPFYLGEKDDVTYAIKPTCWP
						G\LDIIPSCLALHRIETELMGKFD
						EGKLPTDPHLMLRLAIETVAHD
						YDVIVIDSAPNLGIGTINVVCAA
						DVLIVPTPAELFDYTSALQFFD
					, ·	MLRDLLKTVDLKGFEPDVRILL
						TKYSNSNGSQSPWMEEQIRDA
						WGSMVLKNVVRETDEVGKGQI
		l				RMRTVFEQAIDQRSSTDTSLSTP
	ļ					AAPMVDSLIARVGVMARGNAI
						TLPVCGRDVKFTLEVLRGDSVE
						KTSRVWSGNERDQELLTEDAL
						DDLIPSFLLTGQQTPAFGRRVSG
						VIEIADGSRRRKAAALTESDYR
						VLVGELDDEQMAALSRLGNDY
						RPTSAYERGQRYASRLQNEFAG
		į				NISALADAENISRKIITRCINTAK
		i				LPKSVVALFSHPGELSARSGDA
						LQKAFTDKEELLKQQASNLHE
						QKKAGVIFEAEEVITLLTSVLKT
						SSASRTSLSSRHQFAPGATVLY
						KGDKMALNLDRSRVPAYIIRSY
						IRCGRKGFASAGVGGCRGWLN
						YAASEQIVLRVHHMRCEIPHRC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24606	54974	A	24744	1	2624	 MTPQKRGYLCKAGARNLDVLL
						EGCEIFLLIPMGMRSESGVSDA
						DTVSAPRKKHSLEQGIFRAQVE
						ASKPSSVPACIVVGNVSGYQFP
						PGTSYGPDSESELGYREKFMSF
		1				VGTMNEAGNPPFSANYSKAKK
						PKPRMFSLIERMSPSPRVGSQSV
						QSSCFRDIPFEQDQMFPHPSLVL
						RSKEQQPLLTQSIGVNSARKAM
						AKRRHRRRRRVSGVLPIA\VG
						VPS*SVHLPKVLSCSGASWI*QS
						CISRLGRLSPTRLMNVA\GNDP
		1				QGTASMYHGWVQILHIHAEDT
						LL/PFY/LGEKDDVTYAIKPTCW
		İ				PGLDIIPSCLALHRIETELMGKF
		ŀ				DEGKLPTDPHLMLRLAIETVAH
						DYDVIVIDSAPNLGIGTINVVCA
						ADVLIVPTPAELFDYTSALQFFD
						MLRDLLKNVDLKGFEPDVRILL
						TKYSNSNGSQSPWMEEQIRDA
						WASMV*K\NNVRNANEVGKDT
				:		LKGFYSLKERPLGNHFLGPNNR
						PFGKIMKHKTPAFGRRVSGVIEI
						ADGSRRRKAAALTESDYRVLV
i						GELDDEQMAALSRLGNDYRPT
						SAYERGQRYASRLQNEFAGNIS
						ALADAENISRKIITRCINTAKLP
						KSVVALFSHPGELSARSGDALQ
						KAFTDKEELLKQQASNLHEQK
						KAGGFCRFRNHHQTGFSPAGA
						NQRGPLAATLSGPGGEGQSAV
						ARLTGEKKNHPGAQYANRLSP
24607	54975	Α	24745	1	385	MWQAISRLLSEQLGEGEIELRN
				•		ELPGGEVHAAWHLRYAGHDFF
						VKCDERELLPGFTAEADQLELL
						SRSKTVTVPK\AFILGQQIARLH
						QWSDHAVLKIVTTGVL*GAYV
						GAPSIWKCPERITLPGYHQ
24608	54976	Α	24746	1	1209	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24609	54977	Α	24747	1	857	MHACRSELLGEGEIELRNELPG
		1				GEVHAAWHLRYAGHDFFVKC
		ł				DERELLPGFTAEADQLELLSRS
		1				KTVNRALKVWA\VGADRELTV
		1				FLV\MDYLPPRPLDAHSAFILGQ
		1				QIARLHQWSDQPQFGLDFDNA
		į	ļ			LSTTPQPNTWQRRWSTFFAEQR
	1	1		1		IGWQLELAAEKGIAFGNIDAIVE
		İ	Ì			HIQQRLASHQPQPSLLHGDLWS
				1		GNCALGPDGPYIFDPACYWGD
						RECDLAMLPLHTEQPPQIYDGY
						QSVSPLPADFLERQPVYQLYTL
						LNRARLFGGQHLVIAQQSLDRL
24610	54978	Α	24748	1	2094	
24611	54979	Α	24749	1	1209	MVREDLYTYYSCNTSREQQQL
ŕ						ENPMESPLCSKEINQGGVFMLA
						GKASDTLLAGGTMNNLGGEDS
						DTIVENGSIYRLGTDGLQLYSS
						GKTQNLSVNVGGRAEVHAGTL
						ENAVIQGGTVILLSPTSADENFV
						VEEDRAPVELTGSVALLDGAS
						MIIGYGAELQQSTITVQQGAPP
						VKLRFALYGRPGFTKCGWMLG
						YKPAHDKTLLRITRGHDFACYQ
						RTTQLARQRGLKVCSHLIVGLP
						GEGQAECLQTLERVVETGVDGI
						KLHPLHIVKGSIMAKAWEAGR
					<u> </u>	LNGIELEDYTLTAGEMIRHTPPE
						VIYHRISASARRPTLLAPLWCE
						NRWTGMVELDRYLNEHGVQG
		İ				SALGRPWLPPTENSLMKQIRLL
						AQYYVDLMMKLGLVRFSMLL
1				†		A\WPSSFLPLWYKWR
24612	54980	Α	24750	200	310	
24613	54981	Α	24751	1	4398	MCIPVPPIATGDRHFIAVAHIHQ
						NSRRSPVDSDESSTRSAVAAHP
						ATRDFNVRALYTVTSSVSPTKP
	1				}	PLPATFRQGRYRPAEPCVGSIG
						VIPQSSCVLANGSTGAQSAFGDI
		1				IRAGPPSAGLVPVAQATVIETFL
		1				RYQTINERDFWAARNRLLKQP
		1				YGSRILRFTEAYAIAERAAQSV
	.]	1				LRISLADAFNVDGPRPAEDSIFR
[1		1		LLRMPGSAVQRPDRRHNRRCQ
•		1				YRPAMAFPAGSKANKQRQYNG
						GRQFYRKRICWL
	1	1	I	L	l	<u></u>

SEQ ID	1 -	Met	SEQ ID NO:	Nucleotide	i	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24614	54982	ĪΑ	24752	939	2169	RPASPPARDGARWSTGLHSSYS
1						AMRCRNASTCAEDSQLS/DPLA
						R/PSVASGPYRVTSWKLGQNIV
						YCRVKDYWPATLTVNRGRWN
	1					FDTIRSDYYLDDNVAFEAFKAG
		1				AFDLRMENDAK\NWATRYT/G
						DRRVREAITLAFDFEWMNKAL
		İ				FYNAWSRTNSYFQNTEYAARN
		1				SPPPPEL\VLLPPIKKVLPSEVFP\
						QIYQPPVSKGDGYDRDNLLKA
		1				DKLLNEAGWVLKGQQRVNAT
	1	1				TGQPLSFELLLPASSNSQWVLPF
	1				1	QHSLQRLGINMDIRKVDNSQIT
		1				NRMRSRDYDMMPRVWRAMP
		1				WPSSDLQISWSSEYINSTYNAP
	1	İ				GVQSPVIDSLINQIIAAQGNKEK
	Ì				i	FLPLGRALDRVLTWNYYMLPM
]					WYMAEDRLAWWDKFSQPAVR
						PIYSLGIDTWWYDVNKTAKLPS
						ASKQGE
24615	54983	Α	24753	3	450	
24616	54984	Α	24754	2	203	PLPDAARYLRRQCQKPARVHDI
		ŀ				FAAMHTAHRSRPDRPSLPDDAF
		1				SSLRL*FFQPEHYPRQYQHRHL
						CS*ERYQHQALSRTRFRSG*PTP
						SVPETSTGSRYFCGNAHSAPKP
	İ					PRPPITTGRRVFFTTPLILPTRAL
						PASISTPASL
24617	54985	Α	24755	1016	1219	
24618	54986	Α	24756	1695	2021	PDTFSMLPATRFLPPLQFPDLQH
						PEAAQYAHWLPASGCFLFLPPP
						GSGTVFLLPYPSPRLCRSTSETP
		1				LPAITR*HFPVTDRVQK*ITRRR
		<u> </u>				KPRHKPFYPDDLQRDAR
24619	54987	A	24757	1671	1911	OMBAIGA POEDAN VEDGOVONA
24620	54988	Α	24758	513	1104	CWRNCARCERNLVFRCQYGIY
						WCDNAANRSCRTKRDQPRLVE
					1	W*KSACSHHR*LH\PCGHCRQF
						MNELNSGLDLRIHLPGREAHAL
						RDYLPDAFGPKDLEIKTLLMDE
						QDHGYALTGDALSQAAIAAAN
					1	RSHMPYSKSPSGVALECKDGRI\
					1	LVAATLKTPH/INPTQICGQPWS
						MHRRALTSKFGAHGDRRTSAG
						FRRN
24621	54989	Α	24759	131	759	
24622	54990	Α	24760	662	889	LPIKCRAARGVPARCLRWTRW
		1				ALRRILPPLRNRSRAASRWRAS
						PGARK*WMPSLQAVWAAPMR
1						VTRLPAWLRWKC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24623	54991	Α	24761	l l	1518	MQRRDGDIALIYRSKIGPRTSIT
		1				FTARRANPVQRIAARILLRDHLI
		1				CRMSCTAARHANTFDLIQGSG
						ADNFAAIYGGANDRITRADAA
						VAGAGNWHRFGISNGKLSPRSI
	İ		1			TRTSSPARAMTKMPYNPQASG
						GSNLAVSNPELDKLIAARVAAL
		l				RAANPDASASVPVELVTASASY
						PRSLMRCNRILSPGSARGQMIIT
						ALFIYAGNPIACVAALEVLKVF
						EQENLLQKANDLGQKLKDGLL
						AISFWRAKWLKIFVVCGAEIFV
	1					EYVLKNKVIWLLCFANIFLYVV
		İ				RIGIDQWSTVYAFQELKLSKAV
						AIQGFTLFEAGALVGTLLWGW
						LSDLANGRRGLVACIALALIIAT
						LVIPFHMVFHPNEQYAYCVNEL
	1	İ				NSSVDVWELKDPHGNIECVQT
				:		LDMMPENFSDTRWAADIHITPD
	1					GR\HLYACDRTASLITVFSVSED
						GSVLSKEGFQPTETQPRGFNVD
		1				HSGKYLIAAGQKSHHISVYEIV
						GEQGLLHEKGRYAVGQGP\MW
24624	54992	Α	24762	3	486	ACRRALKD\AGVEADEVLEVV
			,_			MVGGSTRVPLVRERVGEFFGRP
						PLTSIDPDKVVAIGAAIQADILV
						GNKPDSEMLLLDVIPLSLGLET
						MGGLVEKVIPRNTTIPVARAQD
	:	İ				FTTFKDGQTAMSAGGAHIRVTF
		l				QVDADGLLSVTAMEKSTGVEA
	:					SIQVKPSYG
24625	54993	В	24763	1	2469	
24626	54994	A	24764	1255	1761	ESFQVRTHGTGEPVNKTRTSQA
						VSHHNQSCQVDQCVPCAVIAG
						NVFPRHNVQGQH*RNCQQTNR
	1	ŀ				SSINDFTKLSNTRPISVHSLGIKQ
						SRPDRHAKASPVVTAVKACEQ
						VIAAVNQIADDHQPDRTKQRN
						AIVVADDLPDLFPVHFFGVDHQ
						QHNHRDKKQPGEDLLCC `
24627	54995	Α	24765	1	3141	
24628	54996	В	24766	1	3969	

SEQ ID . NO:	SEQ ID NO: of peptide sequence	1	09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	54997	A	24767	643	2082	GAHAMDKHFSTTPAEKNLPVL LALIGIWYNNFFDKNDALVMV DDFPRRSGLSWKIVVVPHEYW VGGSTRVPLVRERVGEFFGRPP LTSIDPDKVVAIGAAIQADILVG NKPDSEMLLLDVIPLSLGLETM GGLVEKVIPRNTTIPVARAQDF TTFKDGQTAMSIHVMQGEREL VQDCRSLARFA\CVVFRRYRL/A GAHIRVTFQVDADGLLSVTAM EKSTGVEASIQVKPSYGLTDSEI ASMIKDSMSYAEQDVKARMLA EQKPGQQEAKLALKTFHWQQA IADKRVIGAGITGVTAGTDHRA DVTEVASDFRVQTTNADSALF HVRSAQQHIHQLLHFTTHLLCQ LAGLDHVVFQQIATNPADQVQ AVGFTRTGKDLCHFHGGFTHA EELHKAGVEAGKVTGQAKVEQ MRVQAFYLQQNGADHLRTFW HLNTHRVFHRRGVGGAVGKA ADPAHTVREEGHFVITHTGFRQ FFHAAMDVEQ
24630	54998	Α	24768	1	502	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24631	54999	Α	24769	1	2345	MFMIGALVLALVMTLITFMVL
					1	RRIVIRPLQHAAQRIEKIASGDL
			1			TMNDEPAGRNEIGRLSRHLQQ
	1				}	MQHSLGMTVGTVRQGAEEIYR
						GTSEISAGNADLSSRTEEQAAAI
						EQTAASMEQLTATVKQNADNA
						HHASKLAQEASIKASEGGQTVN
						GVVKTMGAMSTSSKKISEITGR
						HQQGIRTPVTAVKGRCPGPLDE
						GDTDTAYYALRDIFALTPNRKR
						LAIPVSTDNPGKPCKDPSLSKD
						VSNRVPQRVPCYDSPRSSSVNG
						NRLPGKDFHDDERSMGRSGSY
						VLALAYPPKIALSDADSVTVNV
						AGWQGEISREQFNELIAPLVKR
		1				TLLACRRALKD\AGVEADEVLE
		1				VVMVGGSTRVPLVRERVGEFF
i		-				GRPPLTSIDPDKVVAIGAAIQAD
						ILVGNKPDSEMLLLDVIPLSLGL
						ETMGGLVEKVIPRNTTIPVARA
						QDFTTFKDGQTAMSIHVMQGE
						RELVQDCRSLARFACVVFRRYR
		İ				LAGPVDLFLRPWEVDISRRTSL
						DSPLPVQVLEASPKGHYTQLVV
	Į.	l				QPLGWYNEPLTVVMHGDDAP
		ļ				QRGERLFVGLQHARLYNGDER
						ASRDPGFLMARFVTCRPDKTRK
						LRIRQHHVWIEIVSTLEQTIGNT
						PLVKLQRMGPDNGSEVWLKLE
						GNNPAGSVKDRAALSMIVEAE
		İ				KRGEIKPGDVLIEATSGNTGIAL
						AMIAALKSYRMKLLMPDNMSQ
24632	55000	A	24770	313	676	DCQPRSSAFTKPDATTAGHYLA
					070	GSGGRMTQLAIGKPASLGAHLE
						REGGENFTFFSLNADGEKL/CVF
						DANGQ/EHRYDLPGHSGDIWH/
						G/YLPDARPALRMVSRH/GP/WQ
						PAGGMLTPAVVMILPRK
24633	55001	A	24771	1760	2053	FRVHGKNAFSH*IKPAAMLLPS
	1	-				SPVTRSVKPTQCGSAPSKRPNL
						TLLPSITLMPPLPVTDLKLGCIA
						PCSSDCFSFITFSISCATRLSVNW
						LPAAGQ
24634	55002	A	24772	296	732	
24635	55003	A	24773	606	695	
24636	55004	Α	24774	1520	1737	PVHGLCGTRRDAGS\PAIYPFHS
						LML/RFWLR/TCTVPGWPLDDS
						SLAGEAFSAFTTDAAVWSLISA
						TAMLAVQAG
24637	55005	A	24775	1	1049	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Jequence		
24638	55006	Α	24776	248	933	RTTMRAVSRPKYCSRERWL/HY
						DFTVARFDENASCGTFAATSAV
						VLIFSHCLRLLCRVVVLVTRVN
						FQFTEHSTTQRAFWQHAFNRDF
		Ī				NHTLRTASNHLFKGRLFDTTDV
				·		AGVVIVHFVSTLVAGYSNFVSV
	ı					QNDDVITGIYVRSVFRFVLTAQ
						ATSQFSSQTAQSFTGRVNNIPV
ļ						AFYGFWFSCAKYYRHGARWCS
						NGRKIDQRHTHCFFCPCIRTDFA
						VEIGKEEFIT
24639	55007	Α	24777	219	3441	DCQPRSSAFTKPDATTAGHYLA
						GHSGDIWHGYLPDARPGLRYG
						YRVHGPWQPAEGHRFNPAK\LL
						IDPCARQIDGEFKDNPLLHAGH
						NEPDYRDNAAIAPKCVVVVDH
						YDWEDDAPPRTPWGSTIIYEAH
						VKGLTYLHPEIPVEIRGTYKAL
						GHPVMINYLKQLGITALELLPV
						AQFASEPRLQRMGLSNYWGYN
						PVAMFALHPAYACSPETALDEF
	ļ					RDAIKALHKAGIEVILDIVLNHS
		ļ			706	AELDLDGPLFSLRGI
24640	55008	A	24778	454	586	
24641	55009	В	24779	92	1557 2808	
24642 24643	55010 55011	B A	24780 24781	2	1051	WRLRPITPLWRWQKIFFPDA/R
24043	33011	^	24/01	2	1031	YGIRFRNLETGNWYPELLDNVE
	:					PSFVWANDSWIFYYVRKHPIFI
						DLTTEELAASSYHESWVLEDRII
						KDGSYNIDQGVGVRAISGEKTG
						FAYADQISLLALEQSAQAARTI
			<u>†</u>	1		VRDSGDGKVQTLGAVEHSPLY
						TSGRSGWQSMSREEKLDILRRV
						DKVAREADKRVQEVTASLSGV
		1				YELILVAATDGTLAADVRPLGK
						VLLVGQSPNAELSARAKQIAM
						GVDGANEVYNEIRQGQPIGLGE
					1	ASNDTWITTKVRSQLLTSDLVK
						SSNVKVTTENGEVFLMGLVTE
					1	REAKAAADIASRVSGVKRVNR
1						TLFLGYAVMFSELILAPVTPSNS
24644	55012	В	24782	1	2079	
27077	133012	<u></u>	1-1702	1.	1	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24645	55013	IA	24783	1	3214	MGAEKQLIPDGCGVSYIPSSGP
2 10 15	33013	`	2.,,03			LDKWRALLFIQPDYRADPVYR
						QIITITDLRILYDYTENFVLPLSH
						DEVVHGKKSILDRMPGDAWQK
						FANLRAYYGWMWAFPGKKLL
						FMGNEFAQGREWNHDASLDW
						HLLEGGDNWHHGVQRLVRDL
						NLTYRHHKAMHELDFDPYGFE
						WLVVDDKERSVLIFVRRDKEG
						NEIIVASNFTPVPRHDYRFGINO
						PGKWREILNTDSMHYHGSNAG
						NGGTVHSDEIASHGRQHSLSL
24646	55014	Α	24784	1	1079	MKQIKELQRLLGKKTWKMNSS
24040	33014		24704	'		KKPLNMDGPKVDSARALIGRG
				1		WGTHKFYRCDCIKPTIKTNTSS
ľ				1		RGNEKNPDGSHNNIHNSWFRL
				1		LTSQNMCKVPESPAILVDVGQS
					<u> </u>	AETVQSVLTVKVEHPILRIAVA
						GYACQRQSCLLAMRHPQYAIN
						SSKVVKPCSTARQPFQKAYAYP
						PVVRAFPAINVGISGNKLTNDR
i						RYGRTSNKAWGSGDGHTLSEC
						GWAPLAAGDNVGKVSYQTLG
						YYQRILGYIFYPIAW/VMGVPSS
		1				EALQVGSIMA/TKLVSNEFVAM
		ŀ		İ		MDLQK/IASTLSPRAEGIISVFL/
						VSFANFSSIGIIAAAAECRDWFG
						ILAAQTLAVRAALRVVQAERD
						CVLSMEQHQCARGRQR
24647	55015	A	24785	121	258	LTEPRHYCRSIQFFGLTNMGQN
24047	33013	()	24705	121	250	CYQLS*E*FIGTLNHIRRIPQPLS
24648	55016	A	24786	1	470	WGSGDGHTLSECGWAPLAAG
24040	33010	l'`	21700		1,,,	DNVGKVSYQTLGYYQRILGYIF
						YPIAW/VMGVPSSEALQVGSIM
						A/TKLVSNEFVAMMDLQK/IAS
						TLSPRAEGIISVFL/VSFANFSSIG
ŀ						IIAAAAECRDWFGILAAQTLAV
						RAALRVVQAERDCVLSMEQHQ
		İ				CARGRQR
24649	55017	A	24787	1324	1539	GWCGCEVPFKITDGLACLLQSP
24047	33017	l'`	21707	1321		QGHQTFCFT*HNLGLQPFWIPM
ĺ		İ				CLCSVNVFKTRLILTQVEEGCR
		1				AVSMTQ
24650	55018	A	24788	71	403	RTTFIFDLSSGWPFILWQVLSPN
24030	91066		24700	[''	1703	*RCLQVGLFAQIPTGRMGCLHR
1						FLTCRMGVCTDSSPAGWAVTLI
						RMSPIFPASSKSTLSAALLNTSC
ŀ						LHSRDLEPPGAFRSGGPVLL
L	<u> </u>	<u> </u>	<u> </u>	<u> </u>	<u> </u>	LITORDELL FOATROOOF VEL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24651	55019	A	24789	309	1653	SVLMDESDIYRNNRERFFCVFL
	l				LVGVPARGACLGGGGGGCGLCV	
	İ					RYLRSAGWGCGRRFPCVGDN
ļ		ŧ				WSAILAHIGKPEELDTSARNAG
		İ				ALTRREIRDAATLLRLGLAYG
		ł				PGGMSLREVTAWAQLHDVATL
						SDVALLKRLPKCRHWFGILAG
		l				QTLAHIRVLPVIIRAIGFLLSKV
						NGMGKLESFNAVSSLILGQSEN
		1				FIAYKDILGKISRNRMYTMAAT
		l				AMSTVSMSIVGAYMTMLEPKY
		l				VVAALVLNMFSTFIVLSLINPYR
						VDASEENIQMSNLFEGQSFFEM
		l				LGEYILAGFKVAIIVAAMLIGFI
	į					ALIAALNALFATVTGWFGYSIS
		1				FQGILGYIFYPIAWVMGVPSSE
		l		}		ALQVGSIMATKLVSNEFVAMM
						DLQKIASTLSPCAEGIISVLLDSF
		l				ANFSSIGIIARAVKGLNEEQGNE
						GSRFGLKLVYRSTLVS\VLSASI
	ŀ	ļ				AALVLSDHT
24652	55020	Α	24790	754	957	
24653	55021	В	24791	75	3707	
24654	55022	В	24792	75	264	
24655	55023	В	24793	344	461	
24656	55024	A	24794	1	2203	
24657	55025	A	24795	20	198	
24658	55026	Α	24796	1239	1409	RGPPTPFFHVWEWQQRPLSVW
		Ì				EADHV*RRDEGACPRMVARAR
		<u> </u>				NCRPKLLELRAESQ
24659	55027	Α	24797	1325	1564	
24660	55028	Α	24798	323	694	TGYARQKPVPALGWWTNVSAS
		ł				TAPPIILPSITGSRLAVIKWLTVS
						AGKLSALLCASCQKGGGAKWR
	-					MAMANGITNKLAIQCSPPHSTI
						EKFTSQSQHVLA*GVSAVRSM
		<u>L</u>				VSSG*GRRPLPALG
24661	55029	Α	24799	4708	4878	
24662	55030	Α_	24800	20	199	
24663	55031	Α	24801	743	1186	

SEQ ID NO:	SEQ ID NO: of peptide sequence	l .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24664	55032	A	24802	1394	2651	KLMTCRHEGSLGDDPGINGQL WDVNRIDVTAHEGTWERWTV RADEPQAFHIEGVMFQIRNVNG AMPFPEDRGWKDTVWVDGQV ELLVYFGQPSWAHFPFYFNSQT LFSWPQYWAACFGPAPFLPMS REEMDQLGWDSCDIILVTGDA YVDHPSFGMAICGRMLEAQGF RVGIIAQPDWSSKDDFMRLGKP NLFFGVTAGNMDSMINRYTAD RRLRHDDAYTPDNVAGKRPDR ATLVYTQRCKEAWKDVPVILG GIEASLRRTAHYDYWSDTVRRS VLVDSKADMLMFGNGERPLVE VAHRLAMGEPISEIRDVRNTAII VKEALPGWSGVDSTRLDTPGKI DPIPHPYGEDLPCADNKPVAPK KQEAKAVTVQPPRPKPWEKTY VLLPSFEKVKGDKVLYAHASRI LHHETNP\AVPAH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24665	55033	A	24803	1	2126	MQEKQVLAAYVQPSSEGQSRS
24003	33033	^	24003	1	2120	RRARRVSQCHQHSAAAASLAR
			ļ		ļ	TAASATADSAHPPCPRSAARAG
		İ	1			
						ATMHVKKYLLKGLHRLQKGPG
İ						YTYKELLVWYCDNTNTHGPKR
						IICEGPKKKAMWFLLTLLFAAL
						VCWQWGIFIRTYLSWEVSVSLS
						VGFKTMDFPAVTICNASPFKYS
						KIKHLLKDLDELMEAVLERILA
						PELSHANATRNLSFSIVNPTP\L
	•					VLIDERNPHHPMVLDLFGDNH
						NGLTSSSASEKICNAHGCKMA
						MRLCSLNRTQCTFRNFTSATQA
						LTEWYILQATNIFAQVPQQELV
						EMSYPGEQMILACLFGAEPCNY
						RNFTSIFYPHYGNCYIFNWGMT
						EKALPSANPGTEFGLKLILDIGQ
			l			EDYVPFLASTAGVRLMLHEQR
		ĺ				SYPFIRDEGIYPMSGTETSIGVL
						VDKLQRMGEPYSPCTVNGSEV
		ŀ				PVQNFYSDYNTTYSIQACLRSC
		İ				FQDHMIRNCNCGHYLYPLPRG
		:				EKYCNNRDFPDWAHCYSDLQ
						MSVAQRETCIGMCKESCNDTQ
						YKMTISMADWPSEASEDWIFH
]						VLSQERDQSTNITLSRKGIVKLN
						IFFQEFNYRTIEESAANNIVWLL
						SNLGGQFGFWMGGSVLCLIEFG
						EIIIDFVWITIIKLVALAKSLRQR
						RAQASYAGPPPTVAELVEAHT
						NFGFQP\DTAPRSPNIGP\YPGEQ
						ALPNPG/TPAPPNYDSWRLQPL
24666	55034	A	24804	905	1051	
24667	55035	A	24805	1	1257	
24668	55036	Α	24806	1	3116	MPIITLPDGSQRSFDHPVSVAEV
İ						AQSIGAGLAKATLAGKVDGRLI
						ITPKDEEGLEIIRHSCAHLVGHA
						VKQLYPTAKMVIGPVIEEGFYY
						DIFFERPFTPEDMAAIQQRMRE
						LIDKDYDVIKKMTPRAEVIELF
						KLQAAPDRRHAGREGHGLYFH
						EEYVDMCRGPHVPNTRFLKAF
						QLTKISGAYWRGDSKNEQLQRI
İ						YGTAWADKKQLAAYIQRIEEA
						EKRDHRRIGKQLDLFHLQEEAP
						GMVFWHPNGWSVYQ
24669	55037	Α	24807	1924	2790	
24670	55038	Α	24808	1	1185	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24671	55039	A	24809	823	5990	MARPVPLWSSWGLRTSYLPKP LSVNPSRNDGSA/FAGTHPLGV TRGTLAGPAFVRLSLVFLLRRV LAGSSLLPLLMG*QK/PEVLLPD GASEHDYSLRPSDVKRLQNAD LVVWVGPEMEAFMQETGKQIT RSEVSPLCRRVAFNKFNRDPQP RYDWGRLTVLYSGSSLSGGRY VVIIRECHCQLSKPLRLEQNEIR SLLPEASEPIDQAAQEDEAIPQD ELDDKIAGEAGVHEYVVSTGD TLSSILNQYGIDMGD
24672	55040	Α	24810	2305	2550	
24673	55041	Α	24811	665	844	ISFNNEVISYRLSGKNQ*LNLG* EIWLESGAFKKALSPQKHPGLE PSVCTPTPTLVPDR
24674	55042	Α	24812	1	2031	
24675	55043	Α	24813	106	912	HFETVIDQRDFHPVFSHKGIRAT PEGNEEKGQRVADVTEQCGRC RCQPPAATLTQAVKNVHVEDL PQRVGHETGGGDTGDKQVEVG ECAESHTLDALGLRCPEPVMM VRKTVRNMQPGETLLIIADDPA TTRDIPGFCTFMEHELVAKETD GLPYRYLIRKGG**GLIGFDAAF SPHPNPLPRGARGPIALNVAIGL PYLLRNNLNAFAVTSTVAPVSA STASHNPVIQRGDEEDPFQPQR NSDVLADIGVSGACQFNHLHQ AAQVVMG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24676	55044	A	24814	1	1495	MQKLTLTYAETRRLYGKEVYH
24070	33044	Λ.	24014	1	1400	RPSRFIGELPEECVEEVRLRATV
				į		SRPVSHQRMGTPMVENDSGYK
						LGQRVRHAKFGEGTIVNMEGS
						GIAALCHCRVLIATAALMDPDD
						AVELSVTTMMFALCNTRNPRV
						AWDTAPCVPATTAASMAKRAP
			:	·		1
						DMSQSAAPEAGSQKPPRLSCGE
						SLHPLPATKRAYNYCRPVGRIR
						HLCRIRQSMPDATLSRLIRPTTV
						ADPPDKALAPHPAVFTRGDSKF
i i						LSHKPARSPAIRYLLASVWGPV
						GMMAMMLAMTGFVVATRKLR
						RRFGFKSCKNGVLYRISSSSITT
						SPASPIATCRPVLPCWMRMDPM
						LDMMGMQMLMEKYGDQAMA
i						GMDHSQMMGHMFLGHKVIRA
						NHVGDWGTQFGMLFRIAVETF
i i		•				KEDWEAVWDQLDDLNLEGKIV
İ						A*GRDPAAFHIDNGAFTKLSVA
						YALAELVAAVVLDHRRTHPLM
						ADRAAYRGAQAHLFHTLFRQL
						ADKARRAMVNLFTIQTAGFRV
			İ			GQRQFLH
24677	55045	Α	24815	3	707	GCTLGPVKSFSLRVAWLSWPQ
						YFYIIERACSPCCGPLAPGEAPH
		l				VNLFHSTHSENKKQPHNSPYHN
		l				DIQPTLNYLPTNQKKQTQQEQS
						HNITKQGSQLHDVTQVPYQAY
		l .				RSAIVVGLIRRDSVASGIDCRM
						RRERLIRPTGRASDNTEFTGTQ
						NMQNITHSWFVQVMIKATTDA
						WLKGWA*AQRRQPDATPG*RR
						YRTISRQFPATSVLYPAQPAHA
						FTGKYTVYCHRLGQILP
24678	55046	A	24816	2	1340	
24679	55047	Α	24817	1	2457	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24680	55048	A	24818	41	1053	QRNQQAHLSF/QHSRDIYDNQV HLVRNCNLEMPDQEFIVLYGPS DCGKSTTLRMIAGLEEISGGDL LIDGKRMNDVPAKARNIAMVF QNYALYPHMTVYDNMAFGLK MQKIAKEVIDERVNWAAQILG LREYLKRKPGALSGGQRQRVA LGRAIVREAGVFLMDEPLSNLD AKLRVQMRAEISKLHQKLNTT MIYVTHDQTEAMTMATRIVIM KDGIVQQVGAPKTVYNQPANM FVSGFIGSPAMNFIRGTIDGDKF VTETLKLTIPEEKLAVLKTQEV CPDHTLDALGLRCPEPV/HDGA QNRAQYAAWRNVADYRRSG HYPRYSWVLYLYGTRTGC
24681	55049	Α	24819	3	131	
24682	55050	Α	24820	247	492	
24683	55051	A	24821	2	241	YFFQKSSKTVIGGRRLCSVFRY QFSIVLATC*GTFFPSR*STILS* VTQSATSSGREGRSSSRSLNGRF MFSIASMMSW
24684	55052	Α	24822	1	1143	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24685	55053	Α	24823	242	2151	FSRQHVVKMKHCCKNVVILMP
						EPVAEPALNGLRLNLRIVSIVMF
						NFASYLTIGLPLAVLPGYVHDV
						MGFSAFWAGLVISLQYFATLLS
						RPHAGRYADSLGPKKIVVFGLC
		·				GCFLSGLGYLTAGLTAVCLSSA
						VILCRGVQLHPVSDLRIPRGEK
						AGRNRSVDGCCEKAMSVIIVG
						GGMAGATLALAISRLSHGALPV
						HLIEATAPESHAHPGFDGRAIAL
						AAGTCQQLARIGVWQSLADCA
						TAITTVHVVELHNVGQRLFA\L
		i				LRKAPGVTLHCPDRVANVART
						QSHVEVTLESGETLTGRVLVAA
]		DGTHSALATACGVDWQQEPYE
]				QLAVIANVATSVAHEGRAFERF
	1					TQHGPLAMLPMSDGRCSLVWC
		1				HPLERREEVLSWSDEKFCRELQ
		l				SAFGWRLGKITHAGKRSAYPL
						ALTHAARSITHRTVLIQSERLNH
						KALEKRTLTVWDNSFLIYRQSD
		ļ				VSEIRLASLFASFSIRKSKQQAR
						KHTNRRSKDKEKRTRQKRKTT
						SRQRRRRQRMVRVGDEQIAGC
						HCLYSTGSPNRQLREAGIELSDF
		1				VRGNEKARERMKAQYSIAGMT
						SGVVVGTDHAAEAITGFFTKYG
		ŀ				DGGTDINPLYRLNKRQGKQLL
1		ľ				AALACPEHLYKKAPTADLEDD
24686	55054	Α	24824	2	91	
24687	55055	В	24825	1	2346	
24688	55056	Α	24826	1	1713	
24689	55057	Α	24827	1	918	
24690	55058	Α	24828	415	816	
24691	55059	Α	24829	72	296	FCSATFGRPSDQYQNGE*LRWR
						GLGSFACQSPAHENPQWKFVL
						ARRGFARGTAEAHPSQYRHWC
						KIHRLNPGLR
24692	55060	Α	24830	1	620	

SEQ ID		Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
24693	55061	A	24831	1625	2460	GVVNPRQGRVLSMLPAMLLYL
						LFFLIQTSLKSNGGKGKLDPTL
		1				WMWTVNLIYLALAIVLNLWDT
		1				VPVRRLRASFSRKGARIPFVGC
		ļ				RETCELPQIELVKMMLGRELDT
						HALQRAGRTLLSDKPVAAFKN
		ŀ				YGKKGTIAPFDLEVRPGEIVGL
						AGLLGSGRTETAEVIFGIKPADS
						GTALIKGKPQNLRSPHQASVLG
		•				IGFCPEDRKTDGIIAAASVRENII
					:	LALQAQRG\CYVPFPAKNSKRL
						PNALSASLAFAHLQLNNRLNFS
						PAAISKKCCFYVGY
24694	55062	A	24832	1	2034	
24695	55063	Α	24833	3839	3970	
24696	55064	В	24834	694	765	
24697	55065	Α	24835	1778	1982	
24698	55066	Α	24836	1	268	MTCRLLTRLPGQAGVGKRVSG
1						VPDELSPLFSVGSRYLQGGVGL
						VSPTSGANVGLPVISTNNLVKE
						CR\DGKWKMTEKGCLNMFHFK
24699	55067	Α	24837	568	713	
24700	55068	Α	24838	217	1910	AFTFQESVMAFSQAVSGLNAA
ļ						ATNLDVIGNNIANSATYGFKSG
ļ						TASFADMFAGSKVGLGVKVAG
						ITQDFTDGTTTNTGRGLDVAIS
						QNGFFRLVDSNGSVFYSRNGQF
						KLDENRNLVNMQGLQLTGYPA
						TGTPPTIQQGANPTNISIPNTLM
						AAKTTTTASMQINLNSSDPLPT
		1				VTPFSASNADSYNKKGSVTVFD
						SQGNAHDMSVYFVKTGDNNW
						QVYTQDSSDPNSIAKTATTLEF
						NANGTLVDGAM/ANNIATGAIN
						/GLSFLNSMQQNTGANNIVATT
1						QNGYKPGDLVSYQINDDGTVV
						GNYSNEQTQLLGQIVLANFAN
						NEGLASEGDNV\WSATQSSGVA
						LLGTAGTGNFGTLTN/GSIESGI
						AFLVNVPLNPKLKRVNVAGAR
		1				WHPYGWPAFAERACLIEPEGD
		1				RRTHHRPNNQRTYSRKTQRQG
				1		GHKTRQDNWEQKGDGGPRPEP
1						TDPHQHDANRPTVATKKRRKS
1						TQSQRNHRNPKGTHPTEEHGT
						AGTTKPPDTPQTDNWTHKGSL
				1		REGKANRKRNPLRGAEVAGGG
				1		KQRAKAERQTQRGERREDVDE
L						GEEDRSKKTEK
24701	55069	Α	24839	500	653	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24702	55070	Α	24840	445	849	GSPGTAGTEVAVVLKIGGVHD
1						KVIGHYRCSTTNTGRVLDVAI/S
						QNGFFRLVDSNGSVFY/SRNGQ
ŀ						FKLDENRNPATGTPPT/IQQGAN
						PTNISIPNTLMQRKPTAGQIVLA
}				i		NFANHEGLASEATTSGLRRNLL
		l				AWRCM
24703	55071	A	24841	1	453	MKQMDTRGKVINCAFAVSAAF
	ł	İ				ALGDHLGFAAANMNAMIFPMI
ļ						VGKLIGGVTAIGVAMMLVPRY
	}	1				EFIKREISWQSPVFFTPVDKQGG
						LKEAELKTLILEQYHAAGIEPES
						VDSGAIIITGESAKTRNAR*TGR
		1				FKRSGTENL\KLEQYHAAGIEPE
						SVDSGAIIITGESAKTRNARPTV
						MALSQSLGDFVVASAGPHLE
24704	55072	Α	24842	370	670	YSSNIMLRVLSRKALILVPSSSP
		i		ŀ		VKARKPAMLARRGWRSLSRW
						AILSLPGPGR*TGRFKRSGTENL
						\KLEQYHAAGIEPESVDSGAIIIT
	-					GESAKTRNARPAGMALSQSLG
		1				DFVVARAGPHLESVNAGYRTRI
		1				RNHGHSWFPGEIGIRSQFPHTLR
						TGSINGKAWGA
24705	55073	Α	24843	1	1623	
24706	55074	В	24844	1	4986	
24707	55075	В	24845	65	726	
24708	55076	Α	24846	4107	4460	FMSTCGYGGGLSRFPARCLSYQ
İ						PLRSWRETVAASG*RRTGSARA
						NCFCFSGLTRDFSIWRPETK*SA
	}	ł				QTFVVGTWRCGGMGR*IAAVL
						SRYSTVESGVSSTHHRLPLQPDI
		1				PSGRRL
24709	55077	Α	24847	624	895	PLVPAYKQSQRQKPHDYLNRC
						RKSL*QNSTTLHAKNSQ*IRY*
		1				WD/SISE**ELSMTNPQPISY*MG
						KNWKHSL*KLAQDRDALSHRS
						YST
24710	55078	Α	24848	1434	1594	
24711	55079	Α	24849	1294	1488	PAGKQTQQRQQFANDGRGEI
					[WRKLIKPNPRRQKS*CSTDPRQ
						KSTLISQREAVIRRFSEAGNFH
24712	55080	Α	24850	595	879	RRCSTTNIIKKIQQDKKGNRKA
	1					VHRSRFVHGRPITLLFFQLSGRI
						QSHRLPGRAALFAGIPAACDQS
}				*		PSLRQPISSEWSDGR*HRRIFRS*
1]	1	1			SEEF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ļ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24713	55081	A	24851	1009	11478	I IYSLSVSVWAGATVMESPVWM
						PIASKFSIEQTMMQLSFLSRTTSI
						SYSFQPISDSSISNSLVGERSRPR
						AQISSNSSRL*AIPPPLPPMVKD
	1					GRIMHGKPTSAATASASSIVCAI
						PERAVSRPIFFIATSKRRRSSALS
						IASAVAPIMVTPNSSNTP
24714	55082	A	24852	48	209	GLFQVGYAENCRDNNHSFSLH
2 1 / 1 1	33002	1	2 1032	"		SVVIGSSSERVVAKK*PDADHF
						SASFVPVILK
24715	55083	A	24853	1358	3194	PTASFAATLAIGKPVAFDASAE
2 1713	33003	``	24033	1330		ERDTRGFISITIRRPFFGFTANVT
	1					RLETERGGICSRSVPWEVVSKI
						AWAEAPDPPGVTILGLKTQHAT
						/VSERVTSRFTRSHYRQFDLDQ
						AFSAKIFDRYLNLLDYSHKRKT
[ELGDEAAFSAKLDVFYDLYNL
						AQKRRFERYQYALSVLEKPMD
						FTGNDTYNLDRSKAPWPKNEA
						ELNALWDNVFSLAMTAFAREI
						DPHTNYLSPRNTEQFNTEMSLS
						LEGIGAVLQMDDDYTVINSMV
						AGGPAAKSKAISVGDKIVGVG
	1					QTGKPMVDVIGWRLDDVVALI
						KGPKGSKVRLEILPAGKGTKTR
						TVTLTRERIRLEDRAVKMSVKT
						VGKEKVGVLDIPGFYVGLTDD
						VKVQLQKLEKQNVSSVIIDLRS
						NGGGALTEAVSLSGLFIPAGPIV
						QVRDNNGKVREDSDTDGQVFY
						KGPLVVLVDRFSASASEIFARD
						VVVGLCICLSSLDVWLCGLFCR
						CSVDAERTKRQGKKGEESRKR
						RRKGYELSYLLFGWSAVVLLF
						VSDAEQHRASRHGGNACQQHP
						DVRHDEADGYPRQSHQLRLRR
						FRCFRPGRPLRLRRCQHERHDL
						PDDCRQVDRRRNGDWRGDDA
24716	55084	A	24854	100	671	T D D C N Q V D I C C I N C D D I N
24717	55085	A	24855	852	1335	YFCFWTYPLLFPVCMDRKAGK
[<u> </u>				HG\APAGDLYVQVQFNSHPIFE
			1			REGNTLYCEVPINFAMAALGGE
			[IEVPTLDGRVKLKVPGETQTGK
						LFRMRGKGVKSVRGGAQGDLL
						CRVVVETPVGLNERQKQLLQE
						LQESFGGPTGEHNSPRSKSFFD
						GVKKFFDDLTR
24718	55086	A	24856	326	471	PLQTHPAPQQRGHPIYQHGQTV
24/10	133000	Γ	24030	320	7/1	S*SCRRRGWSSISRPVIHHADAQ
1						LMP
	1	<u> </u>	L	.l	<u> </u>	PIAIL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24719	55087	Ā	24857	1095	1283	SCRRRGWSSISRPVIHHADAQL
2.7.13			12.00.		1.202	MP*PQIVYQHLKPPGIYRQQDR
	1					L*IPD*RPVLTIQDVRGP
24720	55088	Α	24858	1	1159	
24721	55089	A	24859	1744	3019	KISPRRPQET/CFQKSTNIAREKT
						LEQEFSVQLFEKIGRRMCLTRE
						GKKLLPHIYELTRVMDTLREAA
						KKESDPDGELRVVSGETLLSYR
						MPQVLQRFRQRAPKVRLSLQSL
						NCYVIRDALLNDEADVGVFYR
						VGNDDALNRRELGEQSLVLVA
						SPQIADVDFTEPGRHNACSFIIN
ŀ						EPQCVFRQIFESTLRQRRITVEN
		Ī				TIELISIESIKRCVAANIGVSYLP
Ì		İ				RFAVAKELECGELIELPFGEQSQ
				,		TITAMCAHHAGKAVSPAMHTFI
			<u> </u>			HALYSCRMRRERLIRATRSCQF
					1	NILHEPRRADKRSASGSLRLQE
		į				FNYQDKPCMEIRIRTYEKTPIR
			ļ			MDKGLTNFGSRWILPELAHKE
			ŀ			KDEETQQRRKSKRDSRREETDT
						KKKTRDKATKREKDKEIEEMA
						SISITSRCEDKLENFSSMKYYTEI
24722	55090	Α	24860	321	575	SYRRYVFPAQ*NRHLQFAVKPP
						APVIVHPTLVDSEHHHSLHDKR
						RQHRPHHSVAHHGSRNSSATA
						TINALLGRVAQNAATRGTR
24723	55091	Α	24861	935	1637	PQPPPPLRREGAGLEKELGGRR
						VRVKGSGCQPWICGKVPVFIPV
						ITEQESFQELI/RRTTTACESLGK
						EYEILMIDDGSSDNSAHMLVEA
		ļ				SQAENSHIVSILLNRNYGQHSAI
						MAGFSHVTGDLIITLDADLQNP
						PEEIPRLVAKADEGYDVVGTVR
						QNRQDSWFRKTASKMINRLIQR
			1			TTGKAMGDYGCM/LRAYRRHI
						VDAMLHCHERSTFIPILANIFAR
						RAIEIPVHHAEHS
24724	55092	Α	24862	1020	1229	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24725	55093	Α	24863	98	1242	MDKFRVQGPTKLQGEVTISGA
						KNAALPNLFAALLAEEPVEIQN
		ŀ				VPKLKDVDTSMKLLSQLGAKV
	•					ERNGSVHIDARDVNVFCAPYD
				1		LVKTMRASIWALGPLVARFGQ
						GQVSLPGGCTIGAHIVMDKVSV
		İ				GATVTIMCAAILTEGATIIEVFY
						FLPLAIGKFNELYPNIKVRIL\EQ
						GTNNCM\ESVLCN\ESDFG\INM\
						NNVTNSSI\DFTPL\VQRRPFVL\
	Ì					ACRRDHPLAKKQLVEWQELVG
	ŀ					YKMIGVRSSSGNRLLIEQQLAD
	Ì					KPWKLDWFYEVRHLSTSLGLV
		ļ.				EAGLGISALPGLAMPHAPYSSII
						GIPLVEPVIRRTLGIIRRKDAVLS
						PAAERFFTLLSDFCPGVPSEDNI
	ł					FESPASQQHEYIGSRQCFLMPSE
	1					TTGTMHV
24726	55094	Α	24864	1	833	MAELGKRSSVAVSWIGKVKTT
		1				AQMVALAWLLWRPPIWVEYA
:						GIALFFVAAVLTLWSMLQYLTL
1						RVQICLISDRFGVIFSKRSKVVK
						NIVDSSRQEYLEEFYARYNVELI
						RAPEGFFYLRPRSTTLIPRSVLS
						ELDMMVGKILCYLYLSPERLA
:						NEGIFTQQELYDELLTLADEAK
						LLKLVNNRSTGSDVDRQKLQE
ŀ						KVRSSLNRLRRLGMVWFMGH
ļ						DSSKFRITESVFRFGADVRAGD
						DPREAQRRLIRDGD/AMPIENHL
						QLNDETEENQPDSGEEE
24727	55095	A	24865	819	1428	SPRGISRGGCRCQPISGDG\RIFA
						GILHGLDNELPLQEEVEGNGLE
ĺ						QEGLPFPIRQSDALDLCFSDILV
						SPNMYIDDGMKRDWPVMKQC
<u>.</u>						ALQRRWRILPDHYGTEPAEKRP
l						GFAPHYCFSYALLVHMMAQQC
		ł				DLEVGDFVWTGGDTHLYSNH
						MDQTHLQLSREPRPLPKLIIKRK
	ŀ					PESIFDYRFEDFEIEGYDPHPGIK
24728	55096	A	24866	2	1445	
24729	55097	A	24867	1	1785	
24/29	2209/	Α	2486/	<u>[1</u>	11/85	<u></u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
<u></u>		<u> </u>				
24730	55098	A	24868	3	654	VRFFKGRGSSWHMV\SQWDYV
						ADRFGTADLLPFTISDMDFATA
İ						PCIIEALNQRLMHGVFGYSRWK
						NDEFLAAIAHWFSTQHYTAIDS
						QTVVYGPSVIYMVSELIRQWSE
						TGEGVVIHTPAYDAFYKAIEGN
	1					QRTVMPVALEKQADGWFCDM
						GKLEAVLAKPECKIMLLCSPQN
						PTGKVWTCDELEIMADLCERH
						GVRVISDEIHMDMVWGEQPHIS
						LE
24731	55099	Α	24869	955	1278	TLTPDFL*QGHFFRFAGIAYQIV
						KQTTVGTTVEITLRRYVVDGNR
			:			IPPRQDRQGDAFFGFATHTQQW
		ŀ	•			HQVFKRQRDIQIVTANTATTVT
						QQAGFTIETRTPLRPHQQQ
24732	55100	Α	24870	1	1707	
24733	55101	A	24871	1	3615	
24734	55102	Α	24872	1	1931	MLLLREHTDTRHDHPRTPTRAT
						RRRRHTEKKRRKKKRPATHVL
		ļ				MTATLIPVSRLLATDQPQLFHE
						SASKPAPHLVSSLGCHCGDASC
						SGRAVADAMQFKTVFPVFVAA
						AMNPEHMAESGHRITRPEFPDY
		1				RELFRESDIKSAVVDSICQCNTP
		l				FNYLFRCFELQCLSRHGHSAYG
		ł				VMQRRHSPMQETRIASLPLQRY
		1				RWHEDSVRNRWFSVMVGPSVR
						VNEWFSAYAMAGMAYSRVST
		<u> </u>				FSGDYLRQPLVSGFHEA
24735	55103	Α	24873	1281	2234	ISTNSILRIFSGFRKLMEV/TRLR
						DTPILTFMNKLDR/DIRDPMELL
						DEVENELK/IGCAPITWPIGCGK
						LFK/GVYHLYKDETYLIQEVRIV
						KGLNNP/DLDAAVGEDLAQQL
						RDE/LELVKGASNEFDKELF/LA
		l				GEITPVFFGTALGNF/GVDHML
						DG\FMAGDRSHVEEAY/PGDIL
		1				G\QGEMMKFTGIPNF/APELFRR
						IRLKDPLKQK/QLLKGLVQLSEE
						GAVQV/FRPISNNDLIVGAVGV
						LQFDVVVARLKSEYNVEAVYE
		1				SVNVATARWVECADAKKFEEF
				1	1	KRKNESQLALDGGDNLAYIATS
						MVNLRLAQERYPGRQFHQTRE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
:	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24736	55104	Α	24874	2	760	AKFEEKFLAVPAEALVYTMKG
						DQKYFPVYANDGKLLPNFIFVA
	1					NIESKDPQQI/DRIQALAGWIAE
	1					Q\IGADVNHATRAGLLSKCDLM
						TNMVFEFTDTQGVMGMHYAR
						HDGEAEDVAVALNEQYQPRFA
]				GDDLPSNPVACALAIADKMDT
						LAGIFGIGQHPKGDKDPFALRR
						AALGVLRIIVEKNLNLDLQTLT
						EEAVRLYGDKLTNANVVDDVI
		1				DFMLGRFRAWYQDEGYTVDTI
1		1				QAVLARRPTRPADFDDWHA
24737	55105	Α	24875	2	391	
24738	55106	Α	24876	2	1785	WLLYRAHVKGE/STEALLPNM
						VATSLAKLPIPKLMRWGATYV
						HFVRPVHTVTLLLGDKVIPATIL
						GIQSDRVIRGHRFMGEPEFTIDN
						ADQYPEILRDSGKVIADYEERK
						AKIKADAEEAARKIGGNADLSE
						TLLEEVASLVEWPVVL/TAKF/E
						EKF/LAISRTALVPYSADNMYQ
						LVNDVQSYPQFLPGCTGSRILES
						TPGQMTAAVDVSKAGISKTFTT
						RNQLTSNQSILMNLVDGPFKKL
						IGGWKFTPLSQEACRIEFHLDFE
						FTNKLIELAFGRVFKELAANMV
ļ					[QAFTVRAKEVYSASMRYIYVIV
						YISYRPPNLLDSHIIGFSSITTHC
						AVTSHHYNSCDMTMTIHSYVT
						PYISVRVSLTQTYAPSPSRNLRI
	1					TDLTIYRAISSEKSIDDTRHIHTI
						ESRCRVCGAHDSRDAERLDRF
						AQTADEIRIADRGFSSRPECIRS
						LAFGEADYIVRVHWRGLRWLT
						AEGMRFDMMGFLRGLDCALIS
						KTRLLSENRRKGRVVQAETLE
						AAGHVLLLTSLPEDEYSAEQVA
						DCYRLRWQIELAFKRLKSLLHL
		1				DALRAKEPELAKVGIFANLLAA
0.4500	55105	<u> </u>	0.4077		2000	FLIDDIIQPSLDFPPRSAGSEKKN
24739	55107	Α	24877	1	2022	
24740	55108	Α	24878	2344	2523	
24741	55109	Α	24879	82	732	

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	,	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24742	55110	Α	24880	828	2184	SGCILPPESLRLWASVARRPRTP
						NQPRQPGPKIGGISPAGSALALA
						CGKRTVLFASLCTRTGGSQLGE
						\PALVLLADQMLALDDQYTFLQ
						RIPRSPYGLAFGPVPHLHYLHQ
						MRLLDIAPIAFRAMSLAFPVNY
						LDAVRTGSICSNCNQRSNDSAM
						GISMNVSTLMRVRAIWVYAAY
		1				MERVLDNLLNNALRYCHSTVE
						TSLLLSGNRATLIVEDDGPGIAP
						ENREHIFEPFVRLDPSRDRSTGG
						CGLGLAIVHSIARGNWATSDHR
						SLTGTQPYARIVVFLVRLIFAFR
						VTQLTLQISFVRFVVVQNAFPG
						CPLGIGIDVHFYYAVAYGVFDF
		İ				LSSRTGTTVFFDDLNTPDFDRC
						FAFMQAVGKGYTDAYLPIVER
		ŀ				RKAMAYGERERNFQLYRRGRY
						VEFNLPSELAATVYQSDPNFAT
						GCFRLYTSCNTFNMHKAATML
						VRAEFSQGYQHQV
24743	55111	Α	24881	679	1628	PDADAKRVDITLCLAFGCDAQF
						IEITFTRLPEVRIHHLQSPASALS
		İ				SCNHLLIRITQFNADITIGHHLP
İ						VNRRVVTVLRDMRHHGDIVHP
						GVRRGVEHYAAMDPGIVKEIM
						EVRLLFLAVFVGNHHSRRDSLP
						VQFVVHTNGDTYTFPGNNIRRD
						IGFKWRIAALMLHHFLIIDPHFA
						VMSDRIETQHDTLACPNFRYKD
						FALIPERCRII/SEYLDRRTDRLK
		1				WCYAAFQQGDQSQASNPMYA
						KAPV*VRWVHGYKALSEKGHL
						VYQRLYQIFQHAEYQTGTNDA
						EISTKLCLFSSSLLLLSSSSLQHSI
						PPNACL
24744	55112	Α	24882	1	1515	
24745	55113	Α	24883	1954	2859	
24746	55114	Α	24884	121	393	SLARWIICNTSEELAANIKRLGD
						VTVRLNAFHAHGVYGEAQSM
						GERHAGRR*AVG*CVKGARGE
						PVDISGAQL/RGCGVNALSGLR
		<u> </u>				FGTDL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	ľ	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24747	55115	IA	24885	1	3706	MDLGANGWQTFRYVVLPNLSS
						ALLAGGMLAFALSFDEIIVTTFT
		-				AEQVDAAVRAADAAFAEWGQ
						TTPKVRAECLLKLADVIEENGQ
						VFAELESSNNTAFARASSNGDL
					1	PTKADLQAQLDSLNKQKDLSA
		1				QDKLVQQDLTDTLATLDKIDRI
		ĺ				KEETVQLRQKVAEAPEKMRQA
						TAALTALSDVDNDEETRKILST
					•	LSLRQLETRVAQALDDLQNAQ
						NDLASYNSQLVSLQTQPERVQ
				_		NAMYNASQQLQQIRSRLDG
24748	55116	Α	24886	2842	5034	NVRLSPAMRCRWRRGSPMPAM
:						FEVGVKTRIIQRSREVILMADHS
		1				KFDAVEPHAVATLSCIKTIISDS
		1				GLPETIAQRYQRAGSSLPEAQP
l		ŀ				DREIEKRGPAIAQAFDAEGKPS
						KASEGWARGCGITVDQAERLT
						TDKGEWLLYRAHVKGESTEAL
						LPNMVATSLAKLPIPKLMRWG
						ASDVHFV\WPVHTVTLL\LGDK
}						VIPATILGIQSDRVIRGHRFMGE
						PEFTIDNADQYPEILRERGKVIA
						DYEERKAKIKADAEEAARKIGG
						NADLSESLLEEVASLVEWPVVL
						TAKFEEKFLAVPAEALVYTMK
						GDQKYFPVYANDGKLLPNFIFV
	ļ					ANIESKDPQQIISGNEKVVRPRL ADAEFFFNTDRKKRLEDNLPRL
		1				QTVLFQQQLGTLRDKTDRIQAL AGWIAEQSGTDAGDQTGYSDE
		1				RWVPDAHSEAPSPVSDMLSAL
						LLLTLVLAGMVRMAARLLMA
						KPPQAVNRGDIGWLTTSPMVIL
						LVMMLAMGTHIPQPVIRIPGGR
		1				FHYSPHQGRTDLPAQRSTWHD
						FCLQAPHLFRRNTVNVNSSSNR
						GEAILAALKTQFPGAVLDEERQ
						TPEQVTITVKINLLPDVVQYLY
						YQHDGWLPVLFGNDERTLNGH
						YAVYYALSMEGAEKCWIVVK
1						ALVDADSREFPSVTPRVPAAV
						WGEREIRDMYGLIPVGLPDQRR
						LVLPDDWPEDMHPLRKDAMD
24749	55117	A	24887	49	531	
24750	55118	A	24888	1619	1968	
24751	55119	A	24889	1	553	
24752	55120	Α	24890	744	970	LAVISEASNQKRGDPVDHAD*I
						AVANMLLNQFQQSYTPMLHDL
						IVRYDGDIHQQKSFGVQLPGKL
						RIVKLIQPVE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24753	55121	Α	24891	52	455	
24754	55122	Α	24892	2988	3614	PVSVLTQKRLSPATSHRSTCRVI
		l	ļ		•	LNQAYVTPSKCWPDVEEISFNF
		1				FHSEDVVRHPVVARIVNAYEA
		1				WEEAEQKTMTPANQHAGGLV
		l				QCDKMLLQRRSREVIELSEQNA
						AQMRHVRGADMAMIFQEPMT
						SLNPVFTVGEQIAESIRLHQNAS
:		ŀ		1		REEAMVEAKRMLDQVRIPEAQ
				1		TILSRYPHQLSGGMRQRVMIA
						MALSCRPAVLIADEPTTALDVT
						IQAQILQLIKVLQKEMSMGVIFI
					1	THDMGVVAEIADRVLVMYQG
						EAVETGTVEQIFHAPQHPYTRA
						LLAAVPQLGAMKGLDYPRRFP
						LISLEHPAKQAPPIEQKTVVDGE
						PVLRVRNLVTRFPLRSGLLNRV
						TREVHAVEKVSFDLWPGETLSL
						VGESGSGKSTTGRALLRLVESQ
						GGEIIFNGQRIDTLSPGKLQALR
			ŀ			RDIQFIFQDPYASLDPRQTIGDSI
		1				IEPLRVHGLLPGKDAAARVAW
						LLERVGLLPEHAWRYPHEFSGG
						QRQRICIARALALNPKVIIADEA
	1					VSALDVSIRGQIINLLLDLQRDF
						GIAYLFISHDMAVVERISHRVA
						VMYLGQIVEIGPRRAVFENPQH
						PYTRK\LLAEDPVAEPSRQRPQR
	1					VLLSDDLPSNIHLRGEEAVVALI
		}				KPGAQIGR\PQSWT
24755	55123	Α	24893	134	595	IRTGRNESYKGDISRYKFVSAR
Ì						ALRNEVRGGRCHKTIRRVSVG
						KKVIGLKFRHLVPVVTPESIVIP
ļ	}					PAPVANDTLVAEVSDAPQAND
	1				}	PTFNNEDLA*FAVLSTPTAILISC
						PGRWGPAAGSECGVRGAHAHR
						ELALRAFSCCCFSSNAISIPN
24756	55124	A	24894	708	1444	TDAPGDEEASLQRAAQAGVGK
						IIVPATEAENFARVLALAENYQ
						PLYAALGLHPGMLEKHSDVSL
						EQLQQALERRPAKVVAVGEIGL
						DLFGDDPQFERQQWLLDEQLIL
						V\KRYELPVILHSRRTHDKLAM
						HLKRHDLPRTGVVYGFSGSLQ
		1				QAERFVQLGYKIGVGGTITYPR
						ASKTRDVIAKLPLASLLLETDA
						PDMPLNGFQGQPNRPEQAARV
						FAVLCELRREPADEIAQAVLNN
				<u> </u>		TYTLVNVP
24757	55125	Α	24895	1460	1777	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24758	55126	Α	24896	381	482	IPGPQSPQSSPDLGWLQC*NP*R LLWKAPRSDK
24759	55127	Α	24897	977	1144	
24760	55128	A	24898	2	280	CTFALRDEYPRMLNVDNKGLR DHIELQPLAV/DASYPGGDFRK NAVYVPMW*SAHCEPEGRFAE TGSRRYGEDFASTAPGFRRAAT RKRVRRKR
24761	55129	В	24899	1	1665	
24762	55130	Α	24900	296	495	
24763	55131	Α	24901	1	5352	
24764	55132	A	24902	187	280	AVDTGLADHRD\EPNMVARLP LSFTASLSIS
24765	55133	A	24903	1775	1981	HIWQNSLIVLFRGCRSAHAKVH RWKN*LPLNLAPLLPRSGSSAPI RPPPSAQARQPMKSTYGVDRR HS
24766	55134	Α	24904	760	1473	
24767	55135	A	24905		2124	MNRMSGGYWQLFAESGETCR ARELEQCLEDFYDQRKESVRW HLDLHTAIRGSLHPQFGVLPQR DIPWDEKFLTWLGAAGLEALV FHQEPGGTFTHFSARHFGALAC TLELGKALPFGQNDLRQFAVTA SAIAALLSGESVGIVRTPPLRYR VVSQITRHSPSFEMHMASDTLN FMPFEKGTLLAQDGEERFTVTH DVEVAKNAITEIKTQFEKEKLA QDRIKRNQ/LDANIQRLNYSLDI ANAAGIKKPVYSN\DGIERKLEI EKAVTDVAELNGELRNRQYLV EQLTKAHVNDVNFTPFKYQLSP SLPVKKDGPGKAIIVILSALIGG MLYCRVCIKRNADSPEVFSGFS DPNVVLIAAVFIIGDGLVRTGV ATVMGTWLVKVAGNSEIKML VLLMLTVAGLGAFMSSTGVVA IFIPVVLSVAMRMQTSPSRLMM PLSFAGLISGMMTLVATPPNLV VNSELLREGYHGFSFFSVTPIGL VVLVLGILYMLVMRFMLKGDT QTPQREGWTRRTFRDLIREYRL TGRARRLAIRPGSPMIGQRLDD LKLRERYGANVIGVERWRFR RVIVNVNGVSEFRARDVSLIAM SAADVDLRQFCSERLLEPDGIA RFSLIPESELIGESVREIGLRTRY GLNVVGLTRTGVALDGSLADE PLLLGDISWLWALESDRYAAM LGRSRSLTYRRGDEHCRASRSP

	of peptide	hod	in USSN	Nucleotide location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, ≔possible nucleotide insertion)
24768	55136	A	24906	834	993	EEIEVARLRDQKSAEGGIQNRE
		l				YWPGRSGHR*SPHAPIIGTQFLQ
				}		ALFQTWDQFVFALSACGRRHP
						TDGRRQNPAVSGHSVAARQPA
						HSQTDEASG
-	55137	Α	24907	3519	5161	
24770	55138	Α	24908	374	1891	KWKLKRPDSNIDHRRVPNLET
						WFSRHDKTRPTSKNPSDYQAG
						DIVSWRLDNGLAHIGVVSDGFA
		ļ				RDGTPLVIHNIGAGAQEEDVLF
						NWRMRQCQHNQHNTATFNGP
İ						RILAEQPPGKEYAANGNGQLYP
		1				MSSLNIKQGSDAHFPDYPLASP
						SNNEIDLLNLISVLWRAKKTVM
						AVVTREENOWOELEKSETKL
		l				AAVVTPPEPVQWQELEKSFTKL RVLDLDIKIDRTEAFNLFIKKFQ
						SVSLLEEYLRSSPYVMDQLKEA
						KIDELDLHRAIVALSEKMKAVD
		l				DNASKKKDEPSLYTSWTLSFTA
						PTSEEAQTVLSGYIDYISTLVVK
		İ	!			ESLENVRNKLEIKTQFEKEKLA
						QDRIKTKNQLDANIQRLNYSLD
						IANAAGIKKPVYSNGQAVKDD
						PDFSISLGADGIERKLEIEKAVT
		l				DVAELNGELRNRQYLV\EPITK
		ļ		1		AHVNDVNFTPFKYQLSPSLPVK
]		KDGPGKAIIVILSALIGGMVAC
						GGVLLRYAMASRKQDAMMAD
24771	55139	A	24909	1111	1895	PLVNFSNSSHPVSNPSIQLSSITL
24//1	33137	ļ'`	24303		1075	PLVENESRARA\VKDQMDAIDH
		l				QTGVRPLVFYSIVLPEIRAIILQS
				-		EGFCQDIVQALVAPLQQEMKL
				•		DPTPIAHRTHGLNPNNLNKYDA
				i		RIAAIDYTLAHDDGISLRNLDQ
						AQVILLGVSRCGKTPTSLYLAM
						QFGIRAANYPFIADDMDNLVLP
						ASLKPLQHKLFGLTIDPERLAAI
						REERRENSRYASLRQCRMEVA
						EVEALYRKNQIPWINSTNYSVE
						EIATKILDIMGLSRRMY
24772	55140	Α	24910	77	434	EKKRGREDKPGTMATFPPATSA
						PQQPPGPEDEDSSLDESDLYSL
						AHSYLGPLIMPMPTSPLTPALV
						TGGGGRKGRTKREAAANTNRP
						SPGGHR/ERKLVTKLQNSERKK
						RGARRLRQSWR
24773	55141	A	24911	31	295	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24774	55142	I A	24912	75	426	 EKKRGREDKPGTMATFPPATSA
24//4	33142	^	24912	1/3	420	PQ\QPPGPGGRDSQPWMDLDLL
						/YSLAHSYLG\GGGRKGHTKRE
						TAANTQPPQPLAGHERKTVDP
						KLQ\NSERKKRGARALRQELEM
						RRRPWTLTPSQ
24775	55143	Α	24913	1	1107	
24776	55144	Α	24914	1	1468	
24777	55145	Α	24915	1	2463	
24778	55146	Α	24916	1	603	
24779	55147	Α	24917	118	1087	IRSKKMTTQTVSGRRYFTKAW
		İ			,	LMEQKSLIALLVLIAIVSTLSPN
						FFTINNLFNILQQTSVNAIMAVG
						MTLVILTSGIDLS/VAASIVGIEV
						NALVAVAAALALGAAIGAVTG
						VIVAKGRVQAFIATLVMMLLL
						RGVTMVYTNGSPVNTGFTENA
						DLFGWFGIGRPLGVPTPVWIMG
						IVFLAAWYMLHHTRLGRYIYA
						LGGNEAATRFSGINVNKIKIIVY
						SLCGQLASLAGIIKVARLSSAQ
						HTAGTGYELDAIAAVVLADDR
						QSGGDFAGGAGRQQKAVITTT
						GHLEYEHEKTGYPGFRCCAKR
						HRQCECDGKRHHRAGGLHA
24780	55148	Α	24918	1	1286	MHEREPGLRAALAGQLEFRVG
		1				LGLGVWTLRLMELLRQRLELV
						VYTASDGQEYEMVSRLNWVN
						YLVLEDVHIILPDQMRGITGCY
						ATHPHERRLHAGTVHRQNIVN
						NPPDCVPRDAGEAEVAPDPSLA
						SLQLQTPELPGLISCHNSIQVSN
l						EARLFELLICQWRDFRKRVIAA
						KSMQGRELEGGQRGQDKATNP
						FRVLALKGSGDPDPHGAGRQG
						SDLLLHPVSDAQVLQHFFQRRA
					<u> </u>	LLSSPWDYHDVVGLCAQEPCG
						YIMELFIIKHQCSLAQTAGSLAP
						EQKKELEMTVRYYISSADLTAE
				}		KFATAIRN/HWHVENKLHWRL
						DV*RQFDKHCSTADSTRILASF
	1					VRHVNAALDVRVVFASGSETT
						HGDGARRYLRCVDVDWRRRT
						AVAQADKSARACHFHYAGYH
						HHEHFADPVSVGAKYHTVFST

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24781	55149	A	24919	417	861	AKGPCPDRNHSSDDKDVIAIDG
						KTLRHSYDKSRR\RGAIHVISAF
	l	ĺ				STMHSLVIGQIKTDEKSNEITAI
		ŀ	:			PELLNMLDIKGKIITTDAMGCQ
						KDIAEKIQKQGVIFSLCPASIIDD
						KKVTDLAVAVQGTLKTGSGGI
						LPPGGALKKISEKL
24782	55150	Α	24920	3	1767	HISIIPDYR*AWKVEHKLSGILL
						LTIFAVISGAESWKDIEDFGKTH
						LDFWKQYGDFENGIPVHDTIAR
	:					VVSCISPAKFHE\CFINWMRDCH
					SSNDKDVIAIDGKTLRHSYDKS	
					RRRGAIHVISAFSTMHSLVIGQI	
						KTDKKSNEITAIPELLNMLDIKG
	İ					KIITTDAMGCQKDIAEKIQKQG
						GDYLFAVKQRNGAVENRAKTQ
						TRIAFGDHFFRRQVSQRIRLFCH
						RLATQFAARLRMFIATGQPMN
						GLRLLLFIVNKQAALLVNKLDA
						QMAAFQQLPGRGRQATVRRE
	i					MDLCAFAERMGLPVSRPGDIQE
						FHAGCRTKFDAGAFRDKHHVA
						CFGDHGLAFVANMQLPFQHHV
						DFVVVQGPREECVLPLHKTGG
						HAAAVVSGHKISQAHAMRLCA
						VETTKLDATGINHFDAAFAGHR
						FESSVDCACSATVSFITFPSVCD
						RFGNWLRSGRCSHAARKNEITL
						TNTRRSLRPKRNTERHDMQTR
						ATVIAAGEKLLPYAETLMSTW
						QAARKEVAHTSRHNEFSIGASA
						SLWDMVMTSAGAVVTTVVVA
						VRPSSFYILCSHKKKRTLHTKR
						EKTVDRRKEYETRAKRSETDT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24783	55151	A	24921	110	1282	SNTPYIIKENKMELKKLMGHISI
						IPDYRQAWKMEHKLSDILLLTI
						CAVISGAEGWEDIQDFGETHLD
						FLKQYGDFENGIPVHDTIARVV
						SCISPAKFHECFINWMRDCHSS
	ļ	ŀ				DDKDVIAIDGKTLRHSYDKST/
						RRRGAIH\VNNAFSTMHSLVIG
						QIKTDKKSNEITAIPELLNMLDI
						KGKIITTDAMGCQKDIAEKIQK
	1					QGGDYLFAVKGNQGRLNKAFE
						EKFPLKELNNPEHDSYAMSEKS
						HGREEIRLHIVCDVPDELIDFTF
						EWKGLKKLCVAVSFRSIIAEQK
						KEPEMTVRYYISSA\DLTAEKFA
						TAIRNHWHVENKLHWRLDVV
						MNEDDCKIRRGNAAELFSGIRH
				1	İ	IAINILTNDKVFKAGLRRKMRK
						AAMDRNYLASVLAGSGLS
24784	55152	В	24922	258	507	
24785	55153	Α	24923	1	3060	MTEIEENMLLPISGQELPIRWLA
						QHGSEKPVTHVSRDGLQALHIA
						RAEELPAVTALAVSHKTSLLDP
						LEIRELHKLVRDTDKVFPNPGN
						SNLGLITAFFEAYLNADYTDRG
						LLTKEWMKGNRVSHITRTASG
						ANACGGNLTDRGDSFVHDLTS
						MARYVATGVLARSMGLVIYSFI
						QHTLTLEEYREITAFLFREFTHA
						AAARYTGEQETLPLSPQWILRQ
						CKEVASLCDGDTFSGEQLNLM
						LQQREWREGFLAER
24786	55154	В	24924	56	967	
24787	55155		24925	1	1776	
24788	55156	Α	24926	1	864	
24789	55157	Α	24927	424	513	
24790	55158	Α	24928	1	5049	
24791	55159	В	24929	1	939	
24792	55160	Α	24930	1	1837	

YVNHFTYSNAIKIFCYQMFLA TQARASGAPLQRLFRAIGGNA WERPFRMVFCSAAFSEESSPFI ETNYEPQLFLGFATDYRFAGV TLRDVEMGYNHDSNGRSDPT RSWNRLYTRLMAENGNWLV! VKPWYVVGNTDDNPDITKYM GYYQLKIGYHLGDAVLSAKG YNWNTGYGGAELGLSYPITKI VRLYTQVYSGYGESLIDYNFN TRVGVGAIGVNVAQAEVLNL SGAKQVLQETFGYQQFRPGQI EIIDTVLSGRDCLVVMPTGGG SLCYQIPALLLNGLTVVVSPLN LKEKSANPFEFTFAAFAHAFH VIQAIGMINTPADGSTHKQAR SWLPYSSLPVLSVSTQDEKAK KMRFMQRSKDSLAK WLSAIL VVIVGLVGLFAVTVIRDYGRA EADRQALLEKGNVLIRALESG RVGMGMRMHHVQQQALLEE AGQPGVLWFAVTDAQGIIILH DPDKVGRALYSPDEMQKLKP ENSRWRLLGKTETTPALEVYF FQPMSAPWRHGMHNMPRCNG KAVPQVDAQQAIFIAVDASDL ATQSGEKRNTLIILYALATVLI ASVLSFFWYRRYLRSRQLLQE	SEQ ID NO:	SEQ ID NO: of peptide sequence	I .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
1 1 1 1 1 1	24793	55161	A	24931		2249	MLPDIARFLPRLANDQIEEFTW YVNHFTYSNAIKIFCYQMFLAS TQARASGAPLQRLFRAIGGNA WERPFRMVFCSAAFSEESSPFR ETNYEPQLFLGFATDYRFAGW TLRDVEMGYNHDSNGRSDPTS RSWNRLYTRLMAENGNWLVE VKPWYVVGNTDDNPDITKYM GYYQLKIGYHLGDAVLSAKGQ YNWNTGYGGAELGLSYPITKH VRLYTQVYSGYGESLIDYNFNQ TRVGVGAIGVNVAQAEVLNLE SGAKQVLQETFGYQQFRPGQE EIIDTVLSGRDCLVVMPTGGGK SLCYQIPALLLNGLTVVVSPLNS LKEKSANPFEFTFAAFAHAFHW VIQAIGMINTPADGSTHKQART SWLPYSSLPVLSVSTQDEKAKK KMRFMQRSKDSLAKWLSAILP VVIVGLVGLFAVTVIRDYGRAS EADRQALLEKGNVLIRALESGS RVGMGMRMHHVQQQALLEEM AGQPGVLWFAVTDAQGIIILHS DPDKVGRALYSPDEMQKLKPE ENSRWRLLGKTETTPALEVYRL FQPMSAPWRHGMHNMPRCNG KAVPQVDAQQAIFIAVDASDLV ATQSGEKRNTLIILYALATVLL ASVLSFFWYRRYLRSRQLLQDE MKRKEKLVALGHLAA/GVAHE IRNPLSSIKGLAKYFAERAPAGG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24794	55162	A	24932	261	2115	TLVFCWHAKLLSHGVFMDIISV
						ALKRHSTKAFDASKKLTPEQAE
						QIKTLLQYSPSSTNSQPWHFIVA
						STEEGKARVAKSAAGNYVFNE
	:					RKMLDASHVVVFCAKTAMDD
İ						VWLKLVVDQEDADGRFATPEA
						KAANDKGRKFFADMHRKDLH
						DDAEWMAKQVYLNVGNFLLG
						VAALGLDAVPIEGFDAAILDAE
						FGLKEKGYTSLVVVPVEDLAGI
				1		EIDHTTSMVMIFGIIFLTAVVVH
						IILHWVVLRTFEKRAIASSRLWL
						QIITQNKLFHRLAFTLQGIIVNIQ
						AVFWLQKGTEAADILTTCAQL
				:		WIMMYALLSVFSLLDVILILGQ\
						KFPA\AFQLPLKG\IFQGIKLMGP
						ILVGILMISLLIGQSPAILISGLGA
						MAAVLMLVFKDPILGL\VAGIQ
						LSANDMLKLGDWLEMPKYGA
						DGAVIDIGLPTVKVRNWDNTIT
						TIPTWSLVSDSFKNWSGMSASG
			!			GRRIKRSISIDVTSIRFLDEDEM
						QRLNKAHLLKPYLTSRHQEINE
						WNRQQGSRESVI\NLRRMP\NIG
						\TFCAYLNK\YLGTH\PRIRKDM
					1	TLMVRQLAPGDNGLPFEIYAFT
						NTVVWLEYESIQADIFDHIFAIV
	•					EEFGLRLHQSPTGNDIRSLAGA
					<u> </u>	FKQ
24795	55163	A	24933	1	757	
24796	55164	Α	24934	8	343	
24797	55165	A	24935	1423	1761	
24798	55166	A	24936	1	851	MKGDTKVINYLNKLLGNELVA
						INQYFLHARMFKNWGLKRLND
						VEYHESIDEMKHADRYIERILFL
						EGLPNLQDLGKLNIGEDVEEML
		ļ				RSDLALELDGAKNLREAIGYAD
		1				SVHDYVSRDMMIEILRDEEGHI
						DWLETELDLIQKMGLQNYLQA
						QIREEGKSIYQEQTTAMKRPDY
						RTLQALDAVIRERGF/EARGTK
				1		AVHYTISRLTAH*ATGKYVRAA
						AVGAYRTAAPDGTRAKTAGTA
						APGGVAGRRVAGR*TNRFDSA
						AAFTGGQRRQSGDVVASCTGS
L	<u></u>		<u> </u>			CVG

SEQ ID	L .		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24799	55167	A	24937	1645	2009	PVFIQFVEGMCGTANEPACLLV
			- 17 4			NQSPAIYFPGKIAAIWQNPRHV
		İ				ESGSM*TL/IDSD*VSPTECLSPR
		1				HHYMPFEEAARGYEIFEKREEE
		1				CRKVILVPGAQSAEAAQKAVS
						GLVNAMPGGTI
24800	55168	Α	24938	1	3936	
24801	55169	Α	24939	1	2240	MRSPMTRNSRNGKHDYKPPPR
						KRRQPPHDHTAPNSTPARNQRG
						RPHRPNRQAGQREQRANNPTP
						NTPSPPPTNDPAEPHQTRHDSR
						QPHNTPPAYAELVAAALLTTDT
						NVTRRRLRYMQPDTKLCHSAG
						KSIYQEQTTAMKRPDYRTLQAL
						DAVIRERGFERAAQKLCITQSA
		1				VSQRIKQLENMFGQPLLVEDET
						RTQERLRRGEVVGAVSIQHQAL
						PSCLVDKLGALDYLFVSSKPFA
1						EKYFPNGVTRSALLKAPVVAFD
						HLDDMHQAFLQQNFDLPPGSV
						PCHIVNSSEAFVQLARQGTTCC
						MIPHLQIEKELASGELIDLTPGL
						FQRRMLYWHRFAPESRMMRK
						VTDALLDYGHKADDIILRITAT
		ŀ				AICGSDLHLYRGKIPQVKHGDI
						FGHEFMGEVVETGKDVKNLQK
						GDRVVIPFVIACGDCFFCRLQQ
				1		YAACENTNAGKGAALNKKQIP
						APAALFGYSHLYGGVPGGQAE
ļ						YVRVPKGNVGPFKVPPLLSDD
						KALFLSDILPTAWQAAKNAQIQ
						QGSSVAVYGAGPVGLLTIACAR
						LLGAEQIFVVDHHPYRLHFAAD
						RYGAIPINFDEDSDPAQSIIEQTA
						GHRGVDAVID\AVGFEAKGSTT
						ETVLTNLKLEGSSGKALRQCIA
						AVKRCGIVSVPGVYAGFIHGFL
						FGDAFDKGLSFKMGQTHVHA
						WLGELLPLIEKGLLKPEEIVTHY
24802	55170	A	24940	687	792	TPTLTRPLRHPVAYRATLK*NR
						PAKVSQLCCTNA
24803	55171	Α	24941	3	510	AGDMVWLCVRTQISY*IPRTHP
						AVGIPRERPLRGADTPQRGASH
						RIRGEGKRK*KRWALLCCPGCN
						I*RHFCHHN*ERYFW\IYWVEST
						GAVKHLQNTGHWLNAGALPV
						RRPRKSVTGSIARQ\VSVPISAR
						ALLACPCGQCAGYATQFVMKS
						ITQRFNVACRTRMYAVG
24804	55172	Α	24942	189	563	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24805	55173	A	24943	1224	1772	GSPSVSLATSSCSIKVQNPGISR VVAGSSAIISNVSPGCILRTVLR TIITGSGQRRPSASSVWSGHTSC VFRTANFSSGMVNLSVSVTNLS PSIVPRIKFIAGDPINPETNIFAG WL/VNGFRRTYLLNNPVFHNH NPRRHGHRFGLIVGHVDHGGV QFLMQLADLCAHLHAQLGIKV
24806	55174	A	24944	1541	3675	LEIIKRKISMRRNWMHSHTRY WRFLDFPTLRGGRWMR*SGEK LNPISTPLRANCLALAQRATRL APMSFTWGMPTSQPDCAVPGR RALPPELLCLQRSSQSRFAGTAI T
24807	55175	В	24945	1	1497	
24808	55176	Α	24946	650	739	
24809	55177	A	24947	4447	4804	RQNDRYSGRERHEITPLCW*IPR IAWALWLRSLRRGPQEKSLHN RVARTISWPYQARHPCRTLLGG HGFFLHIVLSQLPCLSYPLSSFA GLLARSSMFLKLAARFLCCRRS YAASPLSI
24810	55178	Α	24948	1	3774	
24811	55179	Α	24949	431	949	NYAIYHQNHGVNISPGLGKTHN YRQQHCGITYYWRDLMPANK NYKQRWITPGSASFYRSSHKAL IRLLATRLPAFPWGRRSAWRWP VRY*IRVPSSGSYHLEISMRWVI FPQPKALTPSNLHWKTA*LWS* KTLSFVCWIATVLPARFWKLTD PWLAILPCANIYWQASAPD
24812	55180	Α	24950	337	648	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	ł	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24813	55181	Α	24951	1	1430	MFTYPHSKFRVNRFVGQVLLIE
						FSILLDGGRTGLGHSLSVWCISG
	İ					DCSSFLQGKPTSLPLRKPSAIICS
ļ						DSLSGSADLLRNHNFVCTAADT
						FTTDITPEAGDARLIGIAQTVRH
						LEDPMRAWKSIHKSCEILDPIHR
						APVKENTHPARKINISDPAAGA
						VARIRYEGVLSVCRASASAYVS
1						RSRTITVACWYVSNTGVTGATI
1						GVKRQHRHARLFSGDGAPGFG
						TGKRDIRQLRRLVRIQIQRTFFA
						AVLHVPTPWSRAIDAAALLTVI
						DVKVLTACACQACRTLALRVA
	ļ					QVIDRCSNPGDIMSSVARRYWF
ŀ						ISRWSAKRDCAVSYDQDPGQA
İ						QRSCRSCLVAVDRGLWLSAYC
ŀ			*			RSLMTQTLLPDDVPRILPNASSP
ŀ						SRLRCL\HIWVITACWICANSSG
						FIARNL/GLCGWEKRLVKDLTRI
1						QDTGRAKEILGATATLEFR\R*T
1						PTLTRPLRHPVAYRATLK*NRP
1						AKVSQLCCTNA
24814	55182	В	24952	1	1295	AKVSQECTIVA
24815	55183	A	24953	3151	3207	LQWLID*WCL**TFDLRFA
24816	55184	A	24954	2740	3917	NGRSVWAVKLELFRVKPLDIPQ
24010	33104	 ``	24754	2740		QRVGQQHVLLKAFNRRLAMQF
						IQLIATDVTEILPLHVVEDGTDS
						LYCLALSHLRCGRLSLRWRREL
						PRGLGVNVSWCYPFKAQGQCA
						AEITDKAGQREIFGFARVKENA
						GYVIYECYQPDDGDKLIRELPF
						SSLIFARQWFVVGELLQHLPPE
		1				DRITPIVGMLQGVVEKGGELRV
						EVADTNESKELLKFCRKFTVPL
1						RAALRDAGVLANYETPKRPVV
						HVFFIAPGCCYTGYSYSNNNSP
						FYMGIPRLKFPADAPSRSTLKLE
						EAFHVFIPADEWDERLANGMW
						ļ
						AVDLGACPGGWTFQLVKRNM WVYSVDHGSMVQ/SLWITTGT
						1
						CWGTRFHSARGQKFPEVLQGS
04017	55105		24055	1	510	QTLPAATDPPGMQWLASGVLE
24817 24818	55185 55186	A A	24955 24956	510	519 731	
24818	122190	A	24930	مردا	131	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24819	55187	Α	24957	1	1055	MRLCPEKTTKAKRVKAVLAWS
						KEEPWLSEWSEEVLEMGYKKL
						GDFSTAGAILETEIRPSPASAFTL
						DSQPPESNNLSEAKYAPNNIESI
						AKYFAPKSPMFSESPSGTALDS
:						FNRNERRRWPLWPNKSGFGSD
						LTLSKIGISSPDIALERLYEQAR
						KLSGKFRALSELDESARSGVLK
			:			VLCAFAYQDYSRSAASTRKCD
			}			CCDGGGFTEAQVFTNKVSYPW
		1				GKPPYWSKMSRAVRPSDWESW
-			}			TQAREVVRVKCKPCNGKGVIS
į						NSCRCHGKGKVLAKHESDRRG
1						VPVMKACDRCGGRGYARLKFS
1						SV\MRALILLRRLKKTAAYEQL
1						QPLFEELVAECHKQESMADSIL
						SKVTR
24820	55188	Α	24958	361	725	
24821	55189	A	24959	1100	1738	WQRSTLPLLFESAMQTGVNAG
						NPVAW*KSAGYGHIKLYDAKN
						RL\LVQSTALWASQLVQYLSTF
						FRKNLKRPSEFVTLADEIEHVN
						AYLQIEKARFQSRLQVNIAIPQE
						LSQQQLPAFTLQPIVENAIKHGT
						SQLLDTGRVAISARREGQHLML
						EIEDNAGLYQPVTNASGLGMN
						LVDKRLRERFGDDYGISVACEP
					00.40	DSYTRITLRLPWRDEA
24822	55190	A	24960	1	2043	WI WOLLD IN THE CONTROL OF THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE COLUMN TO THE
24823	55191	Α	24961	2164	2487	KLKRIARHNFHGIRDLHLFQQV
			1			NRFSSGFPTTYWLMMADHVHH
						LKTYSTCRIEGGHRFLKDHRHS
						CAAPLTKLFFPRVKNIFSQQGN
l			L	l		LPPMYAITARR*QTHNQHCGD

3295

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		ł		sequence		
24824	55192	A	24962	1128	2574	QVVVVMAMVAKVVIDIVVVV
2,02,				1.23		GVEMIRVKVVAIVVEVVVVEV
		l				VVVMVAWWW****WW*/DMG
		1				EARWWWEYGVWEVEPEIILNA
		l				PQQYLLAGIGDTLAKWYEAVV
		1				LAPQPETLPLTVRLGINNAQAIR
		1				DVLLNSSEQALSDQQNQQLTQS
				ŀ		FCDVVDAIIAGGGMVGGLGDR
					<u> </u>	FTRVAAAHAVHNGLTVLPQTE
						KFLHGTKVAYGILVQSALLGQ
						DDVLAQLTGAYQRFHLPTTLA
]		ELEVDINNQAEIDKVIAHTLRPV
						ESIHYLPVTLTPDTLRAAFKKYS
						QSCFVIRLGGDFIHQLAENHIVI
		i				FVNNHNGTRCQTFQWAISNCH
		İ				TVSLQKFSVRKFLSDLFAIGGR
						RGVRPPTEAPGHQKSDNQAHV
						SVSFDGLDLVFFVVNQWAGVV
		İ			}	FCFDLFSRKRGRFSVWRNKGFG
						RYCQRLANFRDRQAFQLLNQIH
		İ				FTRCKQVDNGGTEFKATHFLAF
						QQCHRTVVFPGRPAIKTRVERL
						DRYRNRMEKSNGQANLRVHG
1		1				GTWEKLVLQLDSAGMVNRVSS
						VGAPLREPIPTEGKGAKEDTWG
		i				NLSSNDPLLEAFGGQFGIAISMH
		l				QTLATRGGKRNARPTTRPRFHR
						GTNPRGIAPHIGTANARYSTGH
		· .				CPVALPFGLRGPPLARENVATA
						TKGHESASRETEVVAVAKLTSR
						VLLICDATHALPIQVLRNDIRVI
						AAGNGCTDDLQLEHLPANTDIR
24825	55193	A	24963	117	335	PGLAPATRRSTPGRFNADAGAD
						*RRAAGTGAGRAKRRGVGIHR
						CLADPFLADCRNAGLRGAALC
1		ł				GTAAASGK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24826	55194	Α	24964	208	1903	YLAIKTFRCLHDFFTDEQLSRA
						VWIYGKRAIAAAQTKLPPAFGL
						PGAKHILFRGHCSESDVQQLAA
						ESGDDRSVVIGVGGGALLDTA
		l				KALARRLGLPFVAVPTIAATCA
						AWTPLSVWYNDAGQALHYEIF
						DDANFMVLVDPEIILNAPQQYL
						LAGIGDTLAKWYEAVVLAPQP
						ETLPLTVRLGINNAQAIRDVLL
1						NSSEQALSDQQNQQLTQSFCDV
						GDAHAGGGMVWGLGDRFTRV
						EAAHAGPNGLTGLPQTEKFLH
						GTKVAYGILV\QSALLGQDDVL
						AQLTGAYQRFHLPTTLAELEVD
						INNQAEIDKVIAHTLRPVESIHY
						LPVTLTPDTLRAAFKKLIQGLIS
	İ					AEQLAQALAEQNGVAWESIDA
						WQIPSSLIAEMPASVALHYAVL
		1				PLRLENDELIVGSEDGIDPVSLA
						ALTRKVGRKVRYVIVLRGQIVT
						GLRHWYARRRGHDPRAMLYN
						AVQHQWLTEQQAGEIWRQYVP
						HQFLFAEILTTLGHINRSAINVL
		1				LLRHERSSLPLGKFLVTEGVISQ
1						ETLDRVLTIQRELQVSMQSLLL
]					KAGLNTEQVAQLESENEGE
24827	55195	В	24965	31	205	
24828	55196	В	24966	69	402	
24829	55197	В	24967	1	721	
24830	55198	Α	24968	1	1703	
24831	55199	Α	24969	2853	2889	LIIGLDLPAVSAIFQRALRFTHV
						GIADGGAYVVQR/HRPDEPRRR
2.4022	155000	ļ	24070	207	4772	CCSE
24832	55200	A	24970	297	473	COVI PRETURI NIVI EL LIDAGII
24833	55201	Α	24971	1179	2148	CQVLRPRTKRLNVLFLHRMQH QWRQRELVHNMCFVIACAKRK
	İ					1 -
	İ					RAPFVQAAEEV\RPGQHDDEFP
						LAIWQTGSGTQSNMNMNEVLA
						NRASELLGGVRGMERKVHPND
						DVNKSQSSNDVFPTAMHVAAL
						LALRKQLIPQLKTLTQTLNEKS
		ŀ				RAFADIVKIGRTHLQDATPLTL
						GQEISGWPASRSGTGSWRYSGG
						YWTKYPSGVCASRSKMNWYST
				1		RASDGGHASDAVRTAQAADRE
						LRAPVRDELNASITSKTRLAEST
						SLAEWVKASPHDDDKPILLPGE
						WEVNTRRERQKQGIPLDAGSW
						QAICDAARQIGMPEETLQAFCQ
L	<u> </u>	<u> </u>		l		Q

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24834	55202	A	24972	166	365	NPFSNSPWETYPVAFTIPLALFM GIYLRYLRPGRIGEVSVIGLV/SP DFRHYLWRLGGRKSDLGTVL
24835	55203	A	24973	83	362	KPPCFLQASVTCWLRLPFASMII LPPCACS*ST*ESIRPAVVGPREP EGITFRRFGRTRVVNRMIFDVL RHLFTVIHPLFQFGVTYITTDDN
24836	55204	Ā	24974	584	853	
24837	55205	A	24975	1	3017	MSQDPFQEREAEKYANPIPSRE FILEHLTKREKPASRDELAVEL HIEGEEQLEGYAPVLAPESGILI MRPTLTMPALTKFVDGTGPVW TGNLFPFLFITIACGAVSGFHALI SSGTTPKMLANEGQACFIGYGG MLMESFVAIMALVSACIIDPGV YFAMNSPMAVLAPAGTADVV ASAAQVVSSWGFSITPDTLNQI ASEVGEQSIISRAGGAPTLAVG MAYILHGALGGMMDVAFWYH FAILFEALFILTAVD
24838	55206	A	24976	3	206	EPDFLPL**RRNHEQNYRDGKC TGSYRSLRTGR*SGQYDHHLRS DPGKSENGRSTGRRRCTGTSVA G
24839	55207	A	24977	3	982	GDAEPFVFLPR\RKDHEYSLDH YQHRFYLRSNRHGKNFGLYRT RMRDEQQWEELIPPRENIMLEG FTLFTDWLVVEERQRGLTSLRQ INRKTREVIGIAFDDPAYVTWIA YNPEPETARLRYGYSSMTTPDT LFELDMDTGERRVLKQTEVPGF YAANYRSEHLWIVARDGVEVP VSLVYHRKHFRKGHNPLLVYG YGSYGASIDADFSFSRLSLLDR GFVYAIVHVRGGGCKINLVVD NGKIVRAEAAQGKTNQGTLCL KGYYGWDFINDTQILTPRLKTP MIRRQRGGKLEPVSWDEALNY VAERLSAIKEKYGPDAIQTTGSS

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
, wo.	sequence	liou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24840	55208	A	24978	1339	3640	SPLSTAGISICGPLLIITDVDLGT
						TVAGIKGMIATGEGTTEWNSV
						MVAMLLTLIPPVVIVLVMQRAF
						VRGLVDMRNKMAGLKLQAVT
		1				KSWDGKTQVIKPLTLDVADGE
						FIVMVGPSGCGKSTLLRMVAG
		İ				LERVTEGDIWINEQRVSELRLIG
		:				YMSRKMGGGKNRREPHIDVKR
	3	}				GIFGLSITPEFSEPLAEDFLSRRQ
		1				YGIRFRNLETGNWYPELLDNVE
			:			PSFVWANDSWIFYYVRKHPVT
		1	ŀ			LLPYQVWRHAIGTPASQDKLIY
						EEKDDTYYVSLHKTTSKHYVVI
						HLASATTSEVRLLDAEMADAE
						PFVFLPRRKDHEYSLDHYQHRF
						YLRSNRHGKNFGLYRTRMRDE
						QQWEELIPPRENIMLEGFTLFTD
						WLVVEERQRGLTSLRQINRKTR
						EVIGIAFDDPAYVTWIAYNPEPE
						TARLRYGYSSMTTPDTLFDLDI
						DTGERRVFKPPE/V/PGFYAA/SY
						RREPRW/IVSADGVKYSFV\APR
				-		QHFRRDAPVLSFLLFSDRAPFR
		1				NNGPRVVLLLGYYHPELLGVT
		1				YSQLSVHEARPKPLPIPHSRHRH
		1				VPTRNEMAQEKERPAGSQSLFR
		1				GLMLIEILSNYPNGCPLAHLSEL
						AGLNKSTVHRLLQGLQSCGYV
		1				TTAPAAGSYRLTTKFIAVGQKA
		1				LSSLNIIHIAAPHLEALNIATGET
		1				INFSSREDDHAILIYKLEPTTGM
		<u> </u>				LRTRAYIGQHMPLYCSAMGKI
24841	55209	Α	24979	1	248	
24842	55210	A	24980	839	1625	PDAFILTTKPAGAIPTSTSITRPIP
		1			1	FCPSFAPCENATPAAET/HQRNT
						RPEWRFTFAFFLFAIFRNAVNT
		1				VTMSIRKGIGVHHNAANLVPW
					}	MTKFFQRFGIAFQSRGAVFHQH
						HHAGYAGQRGKTALAEQRLVF
						GFGINKQMRAVPLIGEVYQFSG
						*CQRQSFTADGFTDSNAFNNIS
		1				GNSTARDKLFIAGYGNRYFHC
						GVNAQIIFMQELGDFRHFVIF/A
						QKGNEYHGDTRISPGRFRRGHH
24942	55211	1_	24001	12529	2964	PLGALRLAASVERSGNGHRA
24843	55211	Α	24981	2538	2864	GSSLINSKVMP/SF*QQNRKFAY
						KRAGVATDEFHAESFVRGRAP
						MKRKL*LFMQLGGDVIQPFDGF
						IAGNTVYRCQQLAVGDAVGKI LHNRYAFRQQSTVIQQQCRHLP
24944	55212	1	24082	1	1201	LINK I AFRQQST V IQQQCRHLP
24844	55212	A_	24982	1	1281	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24845	55213	Α	24983	1	2180	METLNLYADLRSVTGEALLLL
						DEPGVGVDPISRRELWQMVHE
						LAGEGMLILWSTSYLDEAEQCR
						DVLLMNEGELLYQGEPKALTQ
						TMAGRSFLMTSPHEGNRKLLQ
						RALKLPQVSDGMIQGKSNIVGI
						KDTIDSVGHLRTMINTVKSVRP
						SFSVFCGYDDHLLNTMLLGGD
						GAITASANFAPELSVGIYRAWR
						EGDLATAATLNKKLLQLPAIYA
						LETPFVSLIKYSMQCVGLPVET
						YCLPPILEASEEAKDKNRPLVD
						LEHAALQVGKGIIPPPLREYGAS
		l				EVRSVTRAFNHMAAGVKQLAD
						DRTLLMAGVSHDLRTPLTRIRL
						ATEMMSEQDGYLAESINKDIEE
						CNAIIEQFIDYLRTGQEMPMEM
		ĺ				ADLNAVLGEVIAAESGYEREIE
						TALYPGSIEVKMHPLSIKRAVA
						NMVVNAARYGNGWIKVSSGTE
						PNRAWFQVEDDGPGIAPEQRK
		1				HLFQPFVRGDSARTISGTGLGL
						AIVQRIGDNHNGMLELGTSERG
						GLSIRAWLPVPGTRAQGTTKE
						GAREDAVEFFAKLGFVNQGEIT
						TPTTTPIRHFLMIKPVATLDDIL
						HRGDWCAQLQQAWYEHIPLSE
						KMGVRIQQYTGQKFITTMPETG
						NQNPHHTLFAGSLFSLATLTGW GLIWLMLRERHLGGTIILADAHI
						RYSKPISGKPHAVADLGALSGD
						LDRLARGRKARVQMQVEIFGD
24846	55214	A	24984	875	1150	EBREARGICKARVQIMQVEITGB
24847	55215	A	24985	1507	1974	
24848	55216	A	24986	1	1006	
24849	55217	Α	24987	1491	1703	RECRYATNGKNNGCCRSAGSD
						FSSRLRYVR*PFIYTQ*SCGIGH
						CPGTDIRAKIRRTARFIKNYGTY
						CPS
24850	55218	Α	24988	1	1758	
24851	55219	Α	24989	1773	2165	PSRGVEVNSGWNWQATNRRVI
						RNFDDLDQLIIPGTARNAQTCIF
						NFVTVTMTLNNCRRIVQLANQ/
						DYRQPTGTAVAETHGTAEIALL
						ATNFNVAVFIAPLGNQRHNRVL
				1	1	TVWHKFRRVGVLHVGNMASVI
						И
24852	55220	Α	24990	1	141	SPGKIRPVIAAAMKQIKEFMSP
						DSDFFRYMKTPTSATR*YNDCG
						TGQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24853	55221	A	24991	1430	1933	PIIVCMGSLTMPDSACMAPFPPS
				ŀ		AVRRWNSSFPPTFSAHTSSPCA
						CYPRCYRTVKGVL**HHR*W/V
						LISTPGRGAYAASKYALEAWSD
						ALRMELRHSGIKVSLIEPGPIRT
						RFTDNVNQTQSDKPVENPGIAA
						RFTLGPEAVVDKVRHAFISEKP
						KMRYPVTHGDLWR
24854	55222	A	24992	1318	4265	LAIVALFLDLQNHIFSKGFWPS
						DKFYAAKVEEQNQRHQRYHGT
						SYNLEPDIKSSPGGLRDIHTLQ
						WVARRHFGATSLDEMVGFGFL
						TSAERAELNECLHILWRIRFAL
						HLVVSRYDNRLLCVHPTLLTTL
						DIHGEAQIWRREVSSRYGQYPK
		1				AQAAQPDQLMSDYFFRVSLAM
		1				QNKTLLFSLDDTLVNNALQTLN
						KTRPAMVDVIPTDGIVPLYINPQ
						GIAKLLRNETLTSLPKNLEPVFY
						NAAQTLLMPKLDAL
24855	55223	Α	24993	1	454	MGFDDTRDRALFEDSAMSQQP
						TVPDRLRWLLQTFKYQKNIRI*
						STPRKKPMRHSIWNIWGSRRC
						WSNPKRTFMSISVRGNMAAIM
						SFHTSAVMRSHCSILTTIKSRW
		l				ATLNTLILR
24856	55224	Α	24994	65	330	
24857	55225	В	24995	1	1998	
24858	55226	Α	24996	1175	1877	QQVPQKGPGRAKRHLHAAGW
						HWHQRDRCPEEKNVRQ*RRYR
						AVRFIII
24859	55227	В	24997	1	3117	
24860	55228	Α	24998	1888	2412	ARFKHVGDGTVTHRLFRRAAQ
						IVRVV*RAVSHRQHFTGVDIHQ
						HGATRFSLVKGHRIVQFAINQR
						LQAFINTQRQVVRSLAVSRRNI
						FNYATIAVFTHNALTRLPGKPFI
						ETLLNALNPLTEHAARDLEDSP
						VRITAYIEDVDALAQAGADIIAI
						DGTDRPRPVPVETLLARIHH
24861	55229	Α	24999	1	2457	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24862	55230	A	25000		3019	MAALRQSFWSQLFDRVHIGLD FFDASINRIAAWVIGTRNMKKA LLRALLEPTAELRNWKRRAITL RVWHCWKSRNRCRGRRSGKCI ANVTIRQQDHQIETVFINQRLH ERMFPPFVLFAERLSGDEDLGL KAPILEALNDLGYEKPSPIQAEC IPHLLNGRDVLGMAQTGSGKT AAFSLPLLQNLDPELKAPQILVL APTRELAVQVAEAMTDFSKHM RGVNVVALYGGQRYDVQLRA LRQGPQIVVGTPGRLLD
24863	55231	Α	25001	60	545	PSPFTAGASMAVERSTG\HEGE LVLETLVIDGDENT/CAGTRRSL RANA*CLFHPFRAGAEKLAEEV WFCRYPHCGCEQNTTTEEQRR TEWMVTESLADFLDPHDPGKT VEGYPAPKRARNRNRAEDHRQ RLRHRTRKKQTHGQRRRHRPE PAQRQPQTGTTD
24864	55232	Α	25002	1410	1575	
24865	55233	Α	25003	200	691	
24866	55234	Α	25004	1336	2028	
24867	55235	Α	25005	1	345	
24868	55236	A	25006	671	1204	RTSDKLPFCRRSRMENLARFLS TTLKQLRQQRGWSLSRLAEAT GVSKAMLGQIERNESSPTVATL WKIATGLNVPFSTFISPPQSAT/P SLFPYDPQLCFEHLAIQMASGAI SESTPHEKGVIEHVVVIDGQLD LCVDGEWHTLNCGEGVRFAAD VTHIYRNGGEQTVHFHSLIHYP RS
24869	55237	Α	25007	2251	2466	
24870	55238	A	25008	181	626	WQNFDTCGGDMLDKLDAALR FQQEALNLRAQRQEVLAANIA NADTPGYQARDIDFASELKKV MQRGRDATSVVALTMTSTQHI PAQALTPSTAELQYRI\PANPSL DGNTVDMDRERTQFADNSLQY QMSLSALSGQIKGMMNVLQSG N
24871	55239	Α	25009	57	278	
24872	55240	A	25010	171	532	KPVQFAIADGNNQPLALLNIHL PVMMLAIAHGPLLPAFPPELLV LFDDS/PLSSRVLLINAFSGGSDT AMRRAGKGAPAFLLPELPIITV VSPFLPQSSPRRKPIMSKRIPSAV NQRNPRQ
24873	55241	Α	25011	1	648	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i i	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24874	55242	A	25012	158	514	VLQLIKAVWTQRTQKPSWLHP
		l				VDSAPGLQMELPASPVPCTCTP
		1				QPLGGRWDWALWSRGWRLLR
						RLGPHRSPRRS*GSGLWPWPAQ
		l				KGAPIVQRWAKGLLKCHQSGS
		ļ				PGRGGAESERGL
24875	55243	Α	25013	1	693	
24876	55244	Α	25014	119	304	DRRRCAHKITVIPKRMTIAGIRL
						PPQTVNI*LGNFLRALIASTAGI
L						MTAQMVAHSPAPPIP
24877	55245	Α	25015	361	540	
24878	55246	Α	25016	191	446	AKRRLGVYDADCCSIDKTQSLS
		İ				ACTTRNCGAYSKCLCGKYAKP
						VGGIPQPLQESHVR*CCNVMEL
						SPCSDFT**HPPGAVSQPQT
24879	55247	Α	25017	1	2259	
24880	55248	Α	25018	1	1170	
24881	55249	Α	25019	3	1240	
24882	55250	Α	25020	1	1259	MAGPRYPVSVQGAALVQIKRL
						QTFAFSVRWSDGSDTFVRRSW
						DEFRQLKKTLKETFPVEAGLLR
						RSDRVLPKLLGQASLDAPLLGR
						VGRTSRGLARLQLLETYSRRLL
						ATAERVARSPTITGFFAPQPLDL
						EPALPPGSRVILPTPEEQPLSRA
						AGRLSIHSLEAQSLRCLQPFCTQ
		į				DTRDRPFQAQAQESLDVLLRHP
						SGWWLVENEDRQTAWFPAPYL
						EEAAPGQGREGGPSLGSSGPQF
1						CASRAYESSRADELSVPAGARV
						RVLETSDRGWWLCRYAGAGPE
						ELDTSARNAGALTRRREIRDAA
						TLLRLGLAYGPGGMSLREVTA
						WAQLHDVATLSDVALLKRLRN
						AADWFGILAAQTLAVRAAVTG
						CTSGK/QIASCRWNSNQCARGR
						QR*MATTYGI*SSYLSVH*F*AN
						RQQRR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24883	55251	A	25021	803	2061	AKPAARGGILAAPVRDTLFLYF
						RLKKNQRQEGFINTYPWPKKP
						GIFYMCVPIKGANAQDVAAFC
						VTARGRQNCLWESIARGPALIN
						NRGAARGRCNDRVVEVDHGFA
						VLTFNFQGVSPGEFTQTVDHFD
						FAAFRHTSQTTGQLIDHFLFPGT
]			!			NLVDIGFRFAEDDTVFSQRFGF
				1		FDNFCYVQQCFRRDTADVQAN
						TAKGVVTFYDYGFQTYWVKV
						LDPGSEVVAEAGDGASAIDLAH
						RLDIDVILLDLNMKGMSGLDTL
						NALRRDGVTAQIIILTVSDASSD
						VFALIDAGAAGYLLKDSDPEVL
						LEAIRAGAKGSKVFSERVNQYL
						REREMFGAEEDPFSVLTEREL\D
					İ	VLHELAQGLSNKQIASVLNISE
						QTVKVHIRNLLRKLNVRSRVA
						ATILFQSAAFRSRFRRATSDNV
24884	55252	Α	25022	1	1569	
24885	55253	Α	25023	2151	2709	YLREASETARRQFTLP\VILYNF
İ		l				RDLTGQDLTPETVTRLALQNEN
		l				IVGIKDTIDSVGHLRTMINTVKS
		1				VRPSFSVFCGYDDHLLNTMLLG
		ł				GDGAITASANFAPELSVGIYRA
1						WREGDLATAATLNKKLLQLPA
		I				IYALETPFVSLIKYSMQCVGLPV
				1		ETYCLPPILEASEEAKDKVHVL
			1			LTAQGILPV
24886	55254	В	25024	1	1201	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	tocation of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24887	55255	A	25025	1	2209	MIGEIRDGETAEIAIKAAQTGHL
	ļ	ŀ				VLSTLHTNSTCETLVRLQQMG
	ŀ					VARWMLSSALTLVIAQRLVRK
						LCPHCRRQQGEPIHIPDNVWPSP
		l				LPHWQAPGCVHCYHGFYGRTA
	ļ					LFEVLPITPVIRQLISANTDVESL
				}		ETHARQAGMRTLFENGCLAVE
						QGLTTFEELIRVLGMPHGDGAA
						MDREENELGVSCIAVPVFDIHG
						RVPYAVSISLSTSRLKQLYAGLP
						VWCDGSRFFAQKARAİKCKQW
						GYVGAKSRQRWLFYAYDRLRK
					ł	TVVAHVFGERTMATLGRLMSL
						LSPFDVVIWMTDGWPLYESRL
						KGKLHVISKRYTQRIERYNLNL
		1				RQHLARLGRKSLSFSKSVELHD
1						KAAGGVPVLVRELLKAGLLHE
		İ				DVNTVAGFGLSRYTLEPWLNN
						GELDWREGAEKSLDSNVIASFE
						QPFSHHGGTKVLSGNLGRAVM
						KTSAVPVENQVIEAPAVVFESQ
						HDVMPAFEAGLLDRDCVVVVR
						HQGPKANGMPELHKLMPPLGV
						LLDRCFKIALVTDGRLSGASGK
						VPSAIHVTPEAYDGGLLAKVRD
						GDIIRVNGQTGELTLLVDEAEL
						AAREPHISELSAFTFGTGRGYSA
						AFGVKCPFRKRALSDPCYAMR
						EVADFLINKGVDGLFYLGTGGE
				}		FSQMNTAQRMALAEEAVTIVD
						GRVPVLIGVGSPSTDEAVKLAQ
						HAQAYGADGIVAINPYYWKVA
24888	55256	Α	25026	1	3395	MSELPFTIASKRIKYLGIQLTRD
						VKDLFKENYKPLLKEIKEDTNK
						WKYIPCSWLGRINIVKMAILAK
						VIYRFNAIPIKLPMTFFKELEKT
						TLKFIWNQKGAASRSNPEPKNK
		1		1		LEDHITNSTIYRLSTKSCSVSGL
	1					KQTLLAESEALTSYSHRVFSAW
						DFGLCGDVHVRLRQRIILYELK
						GQVPALLLDDGTLLTEGVAIM
						QYLADSVPDRQLLAPVNSISRY
	1				İ	KTIEWLNYIATELHKELKFSFH
						WSTTSHKGLGM
	1		L	I	1	" STISHINGESIVE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24889	55257	A	25027	145	681	FNGATQRIFDARGKPSTYIGSIQ HVNADVDQRATTLQFFIGKDTP AGNAAATQGAGGGNVNLSQN AVGHLFTQLLRIERKTMVKVV
						GQHLAQFARGIDISRACYHMA ARFQCHQGMFVMKTVR*ADA NHIRFHCQQHGLRIVPHGCIPVF FCQRLPLGKAAVTDSYQFQIRC GCSSTKH
24890	55258	Α	25028	696	953	LSGLCSVRRAYPGGRSCPGVP* FVGLVARRPGGG*GSCFVLLHP QRSGTGGCL*QPPFPGAGAGRK IRECIRVGRSAGHAAGGET
24891	55259	A	25029	3	506	RGKPSTYIGSIQHVNADVDQRA TTLQFFIGKDTPAGNAAATQGA GGGNVNLSQNAVGHLFTQLLRI ERKTMVKVVGQHLAQFARGID ISRACYHMAARFQCHQGMFVM KTVR*ADANHIRFHCQQHGLRI VPHGCIPVFFCQRLPLGKAAVT DSYQFQIRCGCSSTKH
24892	55260	Α	25030	291	560	DSTQTQTREGESSTRIT
24893	55261	A	25031	63	404	
24894	55262	A	25032	660	1073	LSHECSFWLTNQLVSPRAQTVS GELQPAYAHGVLCRQYPSAFE KASETPQSSNPCPSSYTIHPTVE NFQTCRNQHLSKGCLINAGRA GPSPGTGGWGELEVAAAARLQ GAWKRHLQSSRPPGQALFASPP *WTGSVQ
24895	55263	A	25033	272	511	GSDTALPSAIAAPLLAAMAWLP PRERGTLAH/QVLEHYQLAQLP VSAL/QMPLHCPPQAIAHHQQL/ EQQALASLQNWGVFHV
24896	55264	Α	25034	1	3300	
24897	55265	A	25035	1117	1376	PPPIGLTYSSSASPLSLFCTVRKP FSRSCW*QTNSNPISSCPELVTP SE*L*PRRSIRLDSLSAVRGFAS RIWLSSAITFPSGSLA
24898	55266	A	25036	207	427	DEHLQQRKKPLFLNYGRTEQA SVK*RISWVQNPERSSLC*GILA A*NPMSVSGL*LT*HCLSARRY ELVCQPK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24899	55267	A	25037	2012	2547	LKFARPKLWGIFRNRQYHHML ER\RSQAIDAPALDRGAALGAL MRLEHPNASAEAALTMLAQLS PAQSGEALHGLLALARHQLAC QPAFIAGFSSHLNQLSEADFINA LPDLRAAMAWLPPRERGTLAH QVLEHYQLAQLPVSALQMPLH CPPQAIAHHQQLEQQALASLQN WGVFHV
24900	55268	A	25038	1	150	DKASRGRSRQQS/CAVLQPLLVI PRQT/WVWSGPPANSSRPGEEG PDC*KEN
24901	55269	A	25039	1	150	DEASRGRNRQQS/CAVLQPLLV IPRQT/WVWSGPPANSSRPAEEE PDC*KEN
24902	55270	A	25040	96	389	PCLSP*LYPSGVHKSTFIYGFYN T*PCTSQSLFWQHSC*GSITSSS YFYLLPEEGTGSNLCCSAASAG DTQANRVWSGPPANSSRPAAE GHDCWKEN
24903	55271	A	25041	1181	1364	VSLGNIKFRQTGAPLGQSFQRK EQSAIFAVLQPPLVIPRQI/WVW SGTPADSSRPAAEGPDC*KEN* RVNRQPTEWEKIFPIYPYDKGLI SRIYKELKQIYKKKIKPPHQKV GKGYVFSKEDIYAANRHMKKC SSSLAIREMQIKTTMRYHLTPA NRGIEIINCPFLDDAASCIDPTYS SNIFPPVNLSWGLPMKPKEVKI KRQKKEHLLKDRDFSFQEPIMA LRTVILEILMEKEMDNSQRECIK DILTKHLVELSILARTFKNTQLP ERAIFQIKQYNSVSCGVSEWQL EEAQVFWAKKEQSLALSILKQ MIKKLDASCAANNPSLKLTYTE CLRVCGNWLAETCLENPAVIM QTYLEKAVEVAGNYDGESSDE LRNGKMKAFLSLARFSDTQYQ RIENYMKSSEFENKQALLKRAK EEVGLLREHKIQTNRCPSGTKL PEEGTVSNLCCSAASTGDTQAN RVWSGTPADSSRPAAEGPDC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24904	55272	A	25042	42	963	VQEGCVRPFCCRSAGVCWRST
						PDSVCLGITSGGCRTAKTAACS
						FLWKLCPRGTPARCQVECSYTR
						CLS\SLLGGVSQAGGTRVRDPL
						EEEVCPLAELERCAGRSVAVFR
						ASRQEHLRLLKLTSQPPLPPGA
						LSQGDGSFIYKPLTGVAAFLSD
						ALPKRRNLERQSGYSTFEELCF
						QKRTWGRDAVRISDLLLGDHS
						HFGKCPISRLERAENMPVVSEL
						SAAGCSTEVNHVFVGIIWQKGK
						CSKYTAPSKSSVPVSSTRGHVHI
						TVQNPTLVPFVGKLNIDSPIGDR
						NIPVLTAGNCKTLGDLRSFPHN
24905	55273	A	25043	98	352	
24906	55274	Α	25044	3	414	PEEALCALAELERCAGRAAALF
						RAERQKRLRLLKLHPQPPLPPG
						ALSQRDGS\LSISP*LGL\RLSFR
						DALPREQFYRLFTDRSDRVPEN
						SVTDVCHIESKMEPLRLGP/SNV
						SIMRPSLVCSFTTLVFISSNAQSS
						LVA
24907	55275	Α	25045	2	1544	STTRGSPPQEARQYRHNQAYA
						YSIQGDGAEDDDERIVRFHTRL
						PVFMLPPTPRKTSWIVAPVLEV
						LARAIRHETEIQGIHTGTEDVPV
						SLFADDMIIYLDNLKDSSRKLL
						EVIRDFSKVSGYKINVHESVAL
						LYTNNDYAENHIKNSTPFTIGY
						VLAFVVMVIAVSCVRLLYAHN
						CTQHTSLKHKCQAWVVTTVDA
		ŀ				AAFMTQATREERRIAYSAPGGY
						FFHILEVLARTIRQEKEIQGIQTG
1						KEEVKVSLFADDMIIYLENLKD
						SSRKLLEVIKEFSKVSGYKINVD
						SRGSPTDLVIYKEEIVFLIVPKA
						KKSKIKVLASDPLCYAATTWR
						MTIIAGLPVEYNDRFIRGIAVFA
						PWRKTPGIYHQSHDPLCYAATT
1						WRMTIIAGLPVEYNDRFIRGIA
						VFAPWRKTPGIYHQSHGACLG
						RRSRTITVVDAQPQVMDMDPT
						CSLFTTGQCLGEPDLLASARRL
						QFFSHQYSIAVLMANARGNSA\
	<u> </u>					LW\DNYGGLIVRADRGSLLLVG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24908	55276	Α	25046	1	1228	MLQSNRTAASHSPSIQPAIQPAI
						QPASQLSIHNLWTQFRHSTKIPV
						EWVNGEWLNGRMPEDHHEWQ
						CNYQNVQQFIDEGNYTSGDNH
						TLRDPHYVEDKGHKYLVFEAN
						TGTENGYQGEESLFNKAYYGG
		1				GTNFFRKESQKLQQSAKKRDA
						ELANGALGIIELNNDYTLKKVM
						KPLITSNTRETVDFNGKHIHVR
						GAKLLFPAIQQPYPPLYFGGSSD
						VAQELAAEQVDLYLTWGEPPE
						LVKDFIEQVRAKAAAHGRKIRF
			ŀ			GIRLHVIVRETNDEAWQAAERL
						ISHLDDETIAKAQAAFARTDSV
						GQQRMAALHNGKRDNLEISPN
						LWAGVGLVRGGAGTALVGDG
						PTVAARINEYAALGIDSFVLSG
						YPHLEEAYRVGELLVPLLDVAI
		1				P\EIPQPQPLNPQGEALRKK
24909	55277	A	25047	1	1075	MTIKGEQAKKQLIAAALAQFGE
						YGMNATTREIAAQAGQNIAAIT
						YYFGSKEDLYLACAQWIADFIG
		•				EQFRPHAEEAERLFAQPQPDRA
ŀ			ļ			AIRELILRACRNMIKLLTQDDT
			l			VNLSKFISREQLSPTAAYHLVH
						EQEAEHRFAMGELPDEVLEICQ
						RLAKLTEMLRGLAELFLNDLSE
1						KTGSHDIVRLHRLILQMNRALG
1						MFEAQSKLWRLASLAQSSGAP
						VTKWATREEREGQLHLWFHCV
		İ				GIRVSDQLERLLWRSIPHIIVTS
		İ				ATLRSLNSFSRLQEMSGLKEKA
:						GDRFVA/SGFP/HLTTANRAKLL
						FPGCALSLPSTTKSSILPKWRPF
						SVSRWRAKNISVCWYCLPADG
						RCSAFSTM
24910	55278	A	25048	530	642	LKTLW/PAVQEPVPHRYHGATV
						FRIMQTIFIKQICRLLL*GTAGHF
						VCGITACHAAEAAHRLTVQPVP
						RILDQRRPAQLHSRQVTAAIRQ
		1				PLHRRQAVRRDAVCQPLLVVC
		1				VFRLMLPQQFPRLRVLHRLAFA
						GQAAFIVIAIVDGNTVAFAQVA
						DVCQPAVAIILPLLLCEFSGQVI
						MITLRPPVKAALLYQPVQRIVT
						EDAVAAVQEPVPHRYHGATVF
						RIMQTIFIKQICRLLL
24911	55279	Α	25049	1075	1479	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence	nou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24912	55280	A	25050	2	588	RWSSLTSSRYGFCRCPESADKE CLHVKSAAGKKATNVGHYAG WLRTYSPHLAAHRVNFFFGINA HMDQRGPGRLMISCNEACGSA CYSNHSYMRNDKHAVRRADSG RAEGGGPRVKSVIKAPVITPKK KPVVSTLKHWVLMLAWVTCSPLTVL\SPDGEMFDVMEKYAFSA WKWKRLRLAISQATFHALLPE KIG
24913	55281	Α	25051	912	1169	
24914	55282	В	25052	1	1665	
24915	55283	Α	25053	43	302	
24916	55284	В	25054	116	1534	
24917	55285	A	25055	1	1017	NVQQFIDEGNYTSGDNHTLRDP HYVEDKGHKYLVFEANTGTEN GYQGEESLFNKAYYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTRETVDFNGKHIHVRGAKLLF PAIQQPYPPLYFGGSSDVAQEL AAEQVDLYLTWGDPADL\VNEI VEQVRAKAAAHGRKIRFGIRLH VIVRETNDEAWQAAERLISHLD DETIAKAQAAF\ARTDSVGQQR MAALHNGKRDNLEISPNLWAG VGLVRGGAGTALVGDGPTVAA RINEYAALGIDSFVLSGYPHLEE AYRVGELLVPLLDVAIP\EIPQP QPLNPQGEALRKK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24918	55286	A	25056	1	2677	MVLNRLLTGLFFGSTPPRCHYD
		-				LIGADPDRTIRSVRLYWRHSLN
						ELISMLPKFNKELTRKLFKLQE
						MLPFLNLHMSPERMLQRMDSE
						KVVTFTTALMTGLSGAMASVL
						LLVMTVVFMLFEVRHVPYKMR
						FALNNPQIHIAGLHRALKGVSH
						YLALKTLLSLWTGVIVWLGLEL
						MGVQFALMWAVLAFLLNYVP
						NIGAVISAVPPMIQVLLFNGVY
				÷		ECILVGALFLVVHMVIGNILEPR
						MMGHRLGMSTMVDGVRVLKD
				ł		GTDLNQTGSFYLAARPYAEKN
				1		GAFIQGVLATFSEADALTRSQR
		-				EQSIALLAKTMGLPAPVIASLLE
						ISQPVSNHPIRTRLTAEVAALQQ
						QTADLFYENRLVPKKVDIRQRI
					,	WQPTRLEGKQFMSLNMFWFLP
		l				THGDGHYLGTEEGSRPVDHGY
		i				LQQIAQAADRLGYTGVLIPTGR
						SCEDAWLVAASMIPVTQR\LKF
						LVALRPSVTSPTVAARQAATLD
						RLSNGRALFNLVTGSDPQELAG
i						DGVFLDHSERYEASAEFTQVW
						RRLLQRETVDFNGKHIHVRGA
						KLLFPAIQQPYPALYFGGSSDV
						AQELAAEQVDLYLTWGEPPEL
						VKEKIEQVRAKAAAHGRKIRFG
						IRLHVIVRETNDEAWQAAERFV
					İ	LSGYPHLEEA/YRVGELLFPLLD
						VAIPE/IPQPQPLNPQGEAVAND
		l				LSPRRTVAASGDQLLAGAVGFS
24919	55287	Α_	25057	114	372	
24920	55288	Α	25058	2270	2760	TSPIPSQIRQYSIA/VGSTGNLGL
				1		SIGIMSARIGFKVTVHMSADAR
						AWKKAKLRSHGVTVVEYEQD
						YGVAVEEGRKAAQSDPNCFFID
		1		1		DENSRTLFLGYSV/AGQRLKAQ
		1				FAQQGRIVDADNPLFVYLPCGV
						GGGPGGVAFGLKLAFGDHVHC
						FFAEPTHSPCMLL
10.400:	55289	В	25059	1	2460	
24921	133209	טן	23037	1,	12400	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
l	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
ĺ				sequence		
24923	55291	Α	25061	332	1349	ILYDSDIYRFSCRSGGHIPAVIY
						HLSAKRRTMENAKMNSLIAQY
						PLVKDLVALKETT/WGFIPATL
						HGVKGLPYVGLTEQDVQDAH\
						ARLSRFAPYLAKAFPGKWLPTG
						GIIES\ELGAL\PAMQKRL\EKEY
			:			QQPISGQLLLKKDSHLPISGSIK
						ARGGIYEVLAHAEKLALEAGLL
ĺ						TLDDDYSKLLSPEFKQFFSQYSI
İ						AVGSTGNLGLSIGIMSARIGFKV
1						TVHMSADARAWKKAKLRSHG
						VTVVEYEQDYGVAVEEGRKAA
						QSDPNCFFIDDENSRTLFLGYSV
						AGQRLK\AQFAQQGRIVDADNP
						LFVYLPCGVGGGPGGVAFGLK
•						LAFGDRVHGLQA
24924	55292	Α	25062	284	1494	RLSLPDMFPPVQQFPVSLHRAF
						YRYDKSAYYALRIAGIASRLNP
						RRFSPSILTPERRCVVTLRNDIR
						RDVLRHNQPRSTASTRQARRR
						MKSSCWHMPTTSKPQALSHTS
		1				NSPTTSISRPNWSYSNVCNRSRT
						MANLSGYNFAYLDEQTKRMIR
						RAILKAVAIPGYQVPFGGREMP
						MPYG\WGTGGIQLTAS\VIGESD
						VLKVIDQGADDTTNAVSIRNFF
						KRVTGVNTTERTDDATVIQTRH
						RIPETPLTEDQIIIFQVPIPEPLRFI
						EPRETETRTMHALEEYGVMQV
-		1				KLYEDIARFGHIATTYAYPVKV
						NGRYVMDPSPIPKFDNPKMDM
						MPALQLFGAGREKRIYAVPPFT
						RVESLDFDDHPFTVQQWDEPC
						AICGSTHSYLDEVVLDDAGNR
}						MFVCSDTDY
24925	55293	Α	25063	1	333	

SEQ ID	SEQ ID NO:		I .		B:	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		02/340,217	sequence	or peptide sequence	detection, (-possible nucleotide insertion)
24926	55294	Α	25064	1368	2451	PAHAAPGGGC/VRSLACRFLWN
1						MRVTGKSYSTLRRVLSAIWNW
						RMMRCALTRALVAFRVR/SSVP
,						LAAVLAIYARENGAGTMFEPE
		l				AAYDEDTSIMNDEEASADNET
}						VMSVIDGDKPDHDDDTHPDDE
						PPQPPRGGRPGLRGLTLRWFSV
		ŀ				AAQRSDILDAVTLSLKVAALAT
						LIALVLGTLAAAALWRRDFFG
1						KNAISLLLLLPIALPGIVTGLALL
						TAFKTINLEPGFFTIVVGHATFC
						VVVVFNNVIARFRRTSWSLVEA
		i				SMDLGANGWQTFRYVVLPNLS
						SALLAGGMLAFALSFDEIIVTTF
			•			TAGHERTLPLWLLNQLGRPRD
						VPVTNVVALLVMLVTTLPILGA
						WWLTAKATYLEKLKV
24927	55295	Α	25065	1	2072	MSEQLTDQVLVERVQKGDQKA
						FNLLVVRYQHKVASLVSRYVP
						SGDVPDVVQEAFIKAYRELTPR
						RPYLLRAFYEWLLDNQLTPHL
1						VVDVTLPGVQVPMEYARDGQI
						VLNIAPRAVGNLELANDEVRFN
						ARFGGIPRQVSVPLAAVLAIYA
						RENGAGTMFEPEAAYDEDTSI
						MNDEEASADNETVMSVIDGDK
			ŀ			PDHDDDTHPDDEPPQPPRGGRP
						ALRVVNVIVAIIFTTKPTAMGIA
						VRFTTAKGETPAIPDVTTTTAVI
			}			GETARASVVAKCIGIIITTAVPIV
ł						VKPFANISPATTSVPRFFGLPAS
						SSHTLIGAIIGIGLTNALLTGSSV
						MDALNLREVTKIFSSLIVSPIVG
ļ						LVIAGGLIFLLRRYWSGTKKRD
						RIHRIPEDRKKKKGKRKPPFWT
						RIALIVSAAGVAFSHGANDGQK
						GIGLVMLVLVGIAPAGFVVNM
						NASGYEITRTRDAVTNFAARES
						EASVIRHSNRFIETINGGRRCSR
						GITLTLHKLR\SAQPLL\AVLNR
				[LEQKKPVGLRYDPQAQSLVCLP
				1		TQTRTGWNLNGFEVGFRPCVR
						LMIYGRSLEAQATASLAAATG
						YDSHIFDLFPASASAQIDTDTAV
						ILLCHDLNRELPVLQAAREAKP
	•					FYLGALGSYRTHTLRLQKLHEL
						GWS\RQETTQIRAPVGIFPKARD
L	<u> </u>					AHTLALSVLAEVASVRLHQEE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24928	55296	A	25066	2	332	AGYSTRTALAIVNKPPSSAIPPT SGLASRKRNPASNDFNGCSRGC GGGGNAGRLNNGSKRSASTPP TIRPTNIAPP\AASPKAPPARWN VSSSGSAPAPRRRANPETQLR
24929	55297	A	25067		3192	MNKKFKYKKSLLAAILSATLLA GCDGGGSGSSSDTPPVDSGTGS LPEVKPDPTPNPEPTPEPTPDPE PTPEPIPDPEP/TPE/PEPTPEPTPD PEPTPEPIPDPEPNQNGAELSYN TEKTLTIRDSVFTYTENADGTIS LQDSNGRKATINLWQIDEANNT VALEGVSADGATKWQYNHNG ELVITGDNATVNNNGKTTVDG KDSTGTEINGNNGKVIQDGDLD VSGGGHGIDITGDSATVDNKGT MTVTDPESM
24930	55298	A	25068	1078	1317	
24931	55299	A	25069	135	361	SCSMLAGLCG*GSSSSVSITTCF ALSASTIA*PREPQQALPV*PLA LPNSSQDGTPMKATSIFSSPLCT RCTRPP
24932	55300	Α	25070	469	841	RWWKQYAAAGFQCMCATAVS NLTRPVAA*TAAIPESKSPVLPN SPASTVRRS*PVSTMPPIRPGDG DHDPRFKAPSGTPHQPPRAGHS L\RRHG*TENCTAA*GQLRTPG DNGTVMSADTITFP
24933	55301	В	25071	1	2922	
24934	55302	Α	25072	1	3849	
24935	55303	Α	25073	1	2280	
24936	55304	A	25074	672	1045	GAVLSINCRTLSPAAASGENVA TPP/VTSNNNTPIINKPKPLALAE ATGVIIAPKPPSSAAPMASPVTF CAVHKPLIRPNIASGVCVCSIVY CSNELTALNQPNSARQTTPKIHI GIRPNRPMARN
24937	55305	A	25075	650	1465	RKPICAADICQLSASAGYSTRT ALAIVSGLASRKRNPASNDFNG CSRGCGGGGNAGRLNNGSKVI NVPAAASP*FQATPPVASNKPPI MGEITDPRRTVNMVIAPACIKF SRPTSAGVAANSAVTPSDAKKP SHKPSK*IASNDGSGSSALRHSP MKPMPQVPRASKITFLRS*RSA STPPTIRPTNIAPPVSVTESPTCQ VECVIIWICTGTATPSKPRDTIE MKRARNTARNAGCVSAGSSGS SAEVGSSNSKTSGSIASARAIET RCC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24938	55306	A	25076	156	3092	LNRRWIANIAKLEAEQARKLK
24730	33300		23070	150	3072	KTEKDSLKDEVLHSLLPRAFSR
			}			FSQTMMWIDTVNGLIMVDCAS
						AKKAEDTLAVLRKSLGSLPVVP
			1			LSMENPIELTLTEWVRSGSAAQ
						GFQLLDEAELKSLLEDGGVIRA
İ						KKQDLTSEEITNHIEAGKVVTK
						LALDWQQRIQFVMCDDGSLKR
						LKFCDELRDQNEDIDREDFAQR
						FDADFILMTGELAALIQNLIEGL
						GGEAQHSGTGSLPEVKPDPTPN
						PEPTPEPTPDPEPTP
24939	55307	D	25077	61	2118	PEPIPEPIPDPEPIP
24939	55308	B A	25078	132	280	RRRAALPERSGPGLDHRRVRH/
24940	122300	A	23078	132	200	GCHVLPTPVTLVKRRKVSRVD
						AQWKSSV
24941	55309	A	25079	1	1449	AQWK33V
24941	55310	В	25080	110	550	
24942	55311	A	25081	657	805	RRRAALPERSGPGLDHRRVRH/
24943	33311	A	23001	037	1803	GCHVLPTPVTLVKRRKVSRVD
						AQWKSSV
24944	55312	A	25082	756	938	AQWK33V
24945	55313	A	25082	781	1257	
24946	55314	В	25083	1	2727	
24947	55315	A	25085	1	2286	
24948	55316	A	25085	1161	2024	VRSAVWPCLQRDPDCHHNRCG
24740	155510		23000	1101	2024	TGGTSRPGPARNPRIEPETCVG\
						RIRNLGVLLYDADAMERAAST
		İ				ENEKDLYQRQLDVFLDPNDPK
						VIEQAIKDGIPLSVIEAAQQSPV
						YKMAMEWKLALPLHPEYRTLP
						MVWYVPPLSPIQSAADAGELGS
						NGILPDVESLRIPVQYLANLLTA
						GDTKPVLRALKRMLAMRHYK
						RAETVDGKVDTRALEEVGLTE
						AQAQEMYRYLAIANYEDRFVV
						PSSHRELAREAFPEKNGCGLPF
						GDGCHGSDTKFNLFNSRRIDAI
						D\KTEPHP
24949	55317	A	25087	664	1065	VAPEPHHGRKQLCREGLGVFH
24747	ا ادودا	^	23001	1007	1003	PTHHITHHLGVSLSLLITIFDAQS
						VHGPDDGCQGLDSVAVDNRLV
						LFHVFSGEAIFMDDLHLLHNGA
						FS*FSSPEQE*F/PRLWRR*CHPL
	<u> </u>	<u> </u>	<u> </u>		<u> </u>	*GFSQSPCSARGRLSPLRSEYIP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24950	55318	Α	25088	434	1663	PAGAFPDIANDFGEKSLLMPGG
1,753						AVINGDF\NNVLPVDLVDPQQV
						QEFVDHAWYRYPNDQVGRHPF
						DGITDPWYNPGDVKGSDTNIQ
						QLNEQERYSWIKAPRWRGNAM
		•				EVGPLARTLIAYHKGDAATVES
						VDRMMSALNLPLSGIQSTLGRI
						LCRAHEAQWAADAIENNYSRN
1						PSQGYSTVPRSPHPKVDPILTKP
						SGPAGPEMTRKQTQREDRPRSR
				•		PSMTTSPTSNLKPSVPAFKGVG
						YLDSLDRIGAADYQPTEQDILR
						TRVKTTGIVETHFTFKNLHFRT
						EQASVHACYCRLFDVEGQRSE
						RKKWIHCFEDVTAIIFCVALSG
						YDQVLHEDETTVLSTRIVDGHR
						YSDHSTIACPAFGTYTLLELKEL
						ALSCSTLLHIPHQRSPAGSFLAD
ļ						NLNCTQAAVIPEFGGF
24951	55319	Α	25089	1	1247	MTFGGDQIFESLVYRSAFRHVY
		1				SRGSPWYLHTNSDLSFVESKKE
						LIAEHILETFPWNLKGLTEIPWG
						DASGESGAVGLGVLAAVHYHP
						QRQSLMEKLALNKDAVVLVIS
						TEGDTDQVIERLESDAVLRYGI
						EDVVTNLDVLERMQPSESLLRA
		İ				VLHTKHLMNPEVLAAARRIVC
		İ				QVVEEIMARLAKEVRQAFSGV
						RDRRRRSFIPLARNFDFKSTLRA
						NLQHWHPQHGKLYIESPRFNSR
		1				IKRQSEQWQLVLLVDQSGSMV
						DSVIHSAVMAACLWQL\PGIRT
						HLVAFDTSVVDLTADVADPVE
						LLMKVQLGGGTNIASAGEYGR
1]			QLIEQPAKSVIILVSDFYEGGSSS
		1	1			LLTHQVKKCVQSGIKVLGLGA
						LDSTATPCYDRDTAQALGNVG
						AQI\AALMPGELGSLAWENLLA

SEQ ID		Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24952	55320	Α	25090	1	1691	MGAYLIRRLLLVIPTLWAIITINF
						FIVQIAPGGPVDQAIAAIEFGNA
						GVLPGAGGEGVRASHAQTGVG
						NISDSNYRGGRGLDPEVIAEITH
						RYGFDKPIHERYFKMLWDYIRF
						DFGDSLFRSASVLTLIKDSLPVS
						ITLGLCWYFVCRACVVTHALSL
			!			WQSASIEVPRTSATAKARKTEP
						GRQTKRKVRSSQTVAARQRIAT
						LPHSMTVCNPGLFHSQMADAL
						AHRYYGFNIAEPWDNGSEAML
						QRDIGLKSESRGLMAGKRCSCL
						KGECGWSLDMQARVSTCFYVR
						LLCGTELEDHRICHDTCDNKCG
						PQRRYVIKFRELCLVNSTVTLV
						DQEYELTVTDPELYDGSSQALA
						GISDLSSIYATCDPGYSCTDRVT
						LVDVRKRQSKTGPYKEIGRYGI
						DRNRSPLNPGFRGKPDLMPSQC
						IACGACACACPANALTIQTDDQ
						QNSRTWQLYMGRCIY/CDGRCE
						EVCPTRAIQ\LTNNFVLTVTNKA
						DLYTRATFHLQRCSRCERPFAP
						QKTIALAAELLAQQQNAPQNR
						EMLWAQASVCPECKQRATLIN
						DDTDVLLVAKEQL
24953	55321	A	25091	1	866	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		•		sequence		
24954	55322	A	25092	338	2388	CVQEPLITVSWRLFAIPAGMKF
						SLFLRCLFCFHLAARYLVLLCQ
						LH/SQGIKVLDFTGVQSGPSCTQ
}		ļ				MLAWFGADVIKIERPGVGDVT
						RHQLRDIPDIDALYFTMLNSNK
						RSIELNTKTAEGKEVMEKLIRE
ŀ]				ADILVENFHPGAIDHMGFTWEH
		1				IQEINPRLIFGSIKGFDECSPYVN
						VKAYENVAQAAGGAASTTGF
						WDGPPLVSAAALGDSNTGMHL
		ŀ				LIGLLAALLHRELNMTVAEIVE
						REEWAGFRARETAALEAVTAP
		ļ				STVIATGGGIILTEFNRHFMQNN
						GIVVYLCAPVSVLEDFRQRTME
						VLRLAFLSSGILEFFTSLSIALVA
						VYFGFSYLGELDFGHYDTGVTL
		l				AAGFLALILAPEFFQPLRDLGTF
		ŀ				YHAKAQAVGAADSLKTFMETP
						LAHPORGEAELASTDPVTIEAE
						ELFITSPEGKTLAGPLNFTLPAG
l						QRAVLVGRSGSGSRYCPYINHR
						NVRLGPIVTDVPWENSGDKSLP
						EATAQVISSGSGATKVEDGLGD
						LNKPVSNQNLVTGIDTPVYNAP
						SAGSAPFGVLADNMRYPILHKL
						KDRLNQTWYQIRIGDRLAYISA
						LDAQPDNGLSVLTYHHILRDEE
						NTRFRHTSTTTSVRAFNNOMA
						WLRDRGYATLSMVQLEGYVK
1						NKINLPARAVVITFDDGLKSVS
	i					RYAYPVLKQYGMKATAFIVTT
24955	55323	Α	25093	1810	2399	STFRNSDDEPYIVRPVAVAWRD
						YVPQPARNGLSNFTGNLEEPAV
						MVNYFLQGDPYQGMVHFTRFF
						HQHPSGLS*WVAGNIFQILCAN
						QPVRKRTRDNVQLLHSRGQRG
						VFRCSRRVLTGLVDRFGRTQTF
ļ						HREAAGEFSGEITGVTDGAWR
						HFRLVLTTQAQRAEEARQQAIS
						GGTEPSAFPDTLPVDTEYVMNF
24956	55324	A	25094	1	577	MPVSTAVATVIKTKQVEFQLQ
						VGVPLYFRLRANPIKTILDNQK
		1				RLDSKGNIKRCRVPLIKEAEQIA
			1			WLQRKLGNAARVEDVHPISER
						PQYFSGDGKSGKIQTVCFEVIG
						CREEPDAEECHGDREQRRVFIR
						EDGFFVTKDLYPNRRA\GIISTT
		1				AVISPSQPPIIAPRVAQSSEPRSA
						AHALARAIRQSTTV
L	<u> </u>		<u> </u>	11	L	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24957	55325	Α	25095	51	407	TCPIRAQRVETLYDRAYREVFG
				-		WNPRGPTGRPNARHTG*SKEG
		İ				GRTEADETDAADSGQRGMEQR
						SEMDRKPTGTTGQRSEKPRRTL
						ECLALHETRIFLPQRKETHSSDA
						HDMPGPPML
24958	55326	Ā	25096	3047	4131	AHSGQMYMPLSSCIRSCILGSV
		İ				VMTCGFPQKSAGQKRQKTADC
						RKRPQLYTKSLSLS/AVKAILAK
		l				GYDPVKWRKKDKVLDAVLKE
						LESGKYSDGDKHAFDQMLHSI
		l				GKQGGDPYLVMADFAAYVEA
						QKQVDVLYRDQEAWTRAAILN
		l				TARCGSSMESKRLDNAALAAGI
			1			SPNYINAHGKPQSISAETKRRLL
				•		DAMHQRTATKVAVTPVPNVM
						VYTSGKKMPMVVEGSGEYSW
						LLTTEEGTQYKGHVTGGKAFN
						LPTKLPEGYHTLTLTQDDQRAH
						CRVIVAPKRCYEPQALLNKPKL
	}	l				WGACVQLYTLRSEKNWGIGDF
	1	į				GDLKAMLVDVAKRGGSFIGLN
						PIHALYPAIRRAPAHTARLLAV
24959	55327	A	25097	524	697	AANWWWKSRRNWR*KMTAM
		\mathbf{I}				LDRIISHIIRTVRMSGSVAVTRAI
						AVPGLLLLIIATA
24960	55328	Α	25098	1	1938	
24961	55329	Α	25099	33	226	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24962	55330	A	25100		2314	MDRAGDGPIRPLSGPTDVYGDT VYLRDPGPNHGKTTNGPLRFVP LLPFGQQVFPRVTPNFWNPVSS AALPARYMQTIENAAVWAQIG DKMVTVGNIRAGQIIAVEPTAA SYYAFNCGFGKAFIDKGHLEPV QGRQKVEDGLGDLYKPLSNQN LVTWKDTPVYNAPSAGSAPFG VLADNLRYPILHKLKDRLNQT WYQIRIGDRLAYISALDAQPDN GLSVLTYHHILRDEENTRFRHT STTTSVRAFNNQMAWARVDRG YATLSMVQLEGYVKNKINNRA TEREELRVSKITVLILGVIAIILG VLFENQNIAFMVGLAFAIAASC NFPIILLSMYWSKLTTRGAMMG GWLGLITAVVLMILGPTIWVQI LGHEKAIFPYEYPALFSITVAFL GIWFFSATDNSAEGAHLRPIRR KKVYRFNYLRSDGPYQGWVSE RLLILRAISPGCSKLVHFPTLFDS LYRTLVKAGEKSGLLAPVLEKL ADYNENRQKIRSKLIQSLIYPC MLTTVAIGVVIILLTAVVPKITE QFVHMKQQLPLSTRILLGLSDT LQRTGPTLLATVFIVAVGFWL WLKRGNNRHRFHAMLLRVALI GPLICAINSARYLRTLSILQSSG VPLLDGMNLSTESLNNLEIRQR LANAAENVRQGNSIHLSLE/QT/ AIFPPMMLYMVASGEKSGQLG TLMGAQSHWECATVNAASRIR
24963	55331	A	25101	20	825	GFPRSRSCYASCTACRTHTYSA EK/QLTRALAKLARATSNEKLS QAFHAHLEETHGQIERIDQVVE SESNLKIKRMKCVAMEGLIEEA NEVIESTEKNEVRDAALIAAAQ KVEHYEIASYGTLATLAEQLGY RKAAKLLKETLEEEKATDIKLT DLAINNSFYKEDIFMNRIEHYH DWLRDAHAMEKQAESMLESM ASRIDNYPELRARIEQHLSETKN QIVQLETILDRNDISRSVIKDSM SKMAALGQSTADTTKNSTHRQ SNTRQGN
24964	55332	A	25102	2	269	GIIGLHLQQQEAVAVAIVRGAT RMLFATLVAVAAGERHYQRW ADC*CAAAG*NVVTVESRTAA ASSHYVGVSRQSHITAVISAPV DVPW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24965	55333	Α	25103	607	727	RQAVYRG**TAGQRANLLLLR
						RHSACALAAGESQSSCPAG
24966	55334	Α	25104	465	574	
24967	55335	Α	25105	3	636	AHTSGFGDKESEKVSPQRKLHA
						MSRTFFWEN\LRCHGNLGSPTY
						AFQISAIDRVNFALVVGLHLRH
						DYRNEKLGRSGRHYQAPVGNR
						SSADGMHRDPCSQLDICCILLG
		1				AGSIYGLYRYIEPQLPDVATLK
				ļ		DVRLQIPMQIYSADGELIAQYG
						EKRRIPVTDGPVMWFPFMIITTC
						VKGNFALNNQFREDMLNYNTE
						GIYGKNVNFENNKQT
24968	55336	Α	25106	434	1205	
24969	55337	Α	25107	368	532	
24970	55338	Α	25108	722	934	
24971	55339	Α	25109	82	420	ARADPHRAHTAGVNGVNARLR
						IFNADTCLRRHAQPCTGGEKDF
		1				RVRFRVFKLGSIRAEGEKLAQL
						QRR*NRRRIQGDRSQPNRD/RA
						VPEDLPSKERMPGRTALTVNSR
						SEWT
24972	55340	Α	25110	618	788	WYPCCRNWRYLAGGDSGFCR
						GERA*R*IASQSGVT*AVATVV
						FPSVLAGLRITGHS
24973	55341	A	25111	186	1198	QKTAVSTSITALTRWGS/SRA/G
						KRGAVLRSRVTRGKYHYPSSW
	ļ	l				RETSSSVQNAPLMRQIKEVFLAI
						RIEQLLTKDEILELYLNKIYLGY
						RAYGVGAAAQVYFGKTVDQL
						TLNEMAVIAGLPKAPSTFNPLY
						SMDRAVARRNVVLSRMLDEG
						YITQQQFDQTRTEAINANYHAP
	İ	1				EIAFSAPYLSEMVRQEMYNRY
						GESAYEDGYRIYTTITRKVQQA
						AQQAVRNNVLDYDMRHAIAA
						RHYVLWKPTANIRARYKAQGI
1						ELPAVVPAGPDNPMGHHAIRL
						AAYGGVYLLHGTNADFGIGMR
						VSSGCIRLRDDDIKTLFSQVTPG
						TKVNIINTPIKVSAEPN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24974	55342	Α	25112	1087	3289	RLRSCGAQKNTFCNAAAGCRW
						RTTLSKMGGLLIAADLQDPVQ
						DMEMWQFHPTGIAC/AGVLVT
						E/GCRGEGGYLLNKHGERFMER
						YAPNAKDLAGRDVVARSIMIEI
						REGRGCDGPWGPHAKLKLDHL
						GKEVLESRLPGILELSRTFAHVD
						PVKEPIPVIPTCHYMMGGIPTKV
						TGQALTVNEKGEDVVVPGLFA
						VGEIACVSVHGANRLGGNSLL
						DLVVFGSARQVLHLQESIAEQG
						ALRDASESDVEASLDRLNRWN
						NNRNGEDPVAIRKALQECMQH
						NFSVFREGDAMAKGLEQLKVI
						RERLKNARLDATSSEFNTQRVE
		İ				CLELHNLMETAYATAVYATFR
	}					TERRGAHSR\FDFPDRDDENWL
	ŀ			1		CHSLYLPESESMTRRSDYTLEA
						DEGRDMMLLDALIQLKEKDPS
						LSFRRSCREGVCGSDGLNMNG
			:			KNGLACITPISALNQPGKKIVIR
						PLPGLPVIRDLVVDMGQFYAQ
						YEKIKPYLLNNGQNPPAREHLQ
	İ					MPEQREKLDGLYECILCACCST
						SCPSFWWNPDKFIGPGSLLAGY
						RSGLIAGIPKCSARYLSGEKRSV
						AWQQIAAEIGISIDAQFNESLKG
						ISRDESLRRILQHGGKEGDFNSQ
						ERAQLAYRKNLLYVHSLRELT
						VNAVLPGIRSLLADLRAQQISV
						GLASVSLNAPTILAALELREFFT
						FCADASQLKNSKPGPGIFLAGC
24975	55343	Α	25113	47	630	VGWETSFAQIRSRFLRIRFCRAY
						SSTLFVSAANPTT*G/QMVF*CG
						NGAQHKNILLLYRSIHGGFHFQ
						RGHHINTFNKRWRR**NRT*YQ
						RDVRATTGSGGGDRKTHLAGA
						VIRDITHRIERFTRRAGGNHNV
						QMFQVVDGRKCATAISTVLDM
		1				FGDHFGRQKIASCQRWLAKTIR
						TVQQPICKNIVRKPGCHSLA
24976	55344	Α	25114	731	829	
				·	J	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24977	55345	A	25115	2	3744	RLPFKYPVKVKPLPAPFSKILLN
						SPAIPQLPITMNGSDFSLPEGPM
	1					NRWENIQLTPETRLAPRAYFFS
	1					YDSVAQARTFARETSSLFLPLS
						GQW\NFHFFDHPLQVPEAFTSE
		İ				LMADWGHITVPAMWQMEGHG
			ŀ			KLQYTDEGFPFPIDVPFVPSDNP
	Ì					TGAYQRIFTLSDGWQGKQTLIK
			ł			FDGVETYFEVYVNGQYVGFSK
						GSRLTAEFDISAMVKTGDNLLC
			ŀ			VRVMQWADSTYEKDQDMW.W
ŀ						SAGIFRDVYLVGKHLT
24978	55346	A	25116	1	2643	SAGII RBVI EVGRILLI
24979	55347	A	25117	2499	2935	RSGGVACNCAVSASCMLVTVL
24313	33347	Ι^	23117	2499	2733	PWLSLPV/SVFGLWDGFFASLG
						VTALLAATPA\W*GVKT*CRPG
						RSP/SLTVRLGSVISPMIGGLLLA
						TGGVALLPLLSLPALPPPPQPRE
						HPLKSLLAGFRFLLASPLVGGIA
						LLGGLLTMASAVRV
24980	55348	A	25118	676	1035	LEGGLETMASAVKV
24981	55349	A	25119	1	1842	
24982	55350	A	25120	1	1341	
24983	55351	A	25121	563	1360	PSVLRCRNWVI*SPYCISLPSRV
12.703						SSLPFLVWIPVAR/CTAIGASRE
						AMLGVLVEPMLLLGLWVAAQ
						VAGSTNISNITDTVYHWPLSQSI
						PLVLALCACAFATFIEMGKLPF
						DLAEAEQELQEGPLSEYSGSGF
						GVMKWGISLKQLVVLQMFVG
						VFIPWGQMETFTAGGLLLALVI
						AIVKLVVGVLVIALFENSMARL
		1				RLDITPRDMIRIELDSWEQNGE
		1				AIKRTGQPLLLSLLVRNLLDNA
		ł				VRYSPQGSVVDVTLNADNFIVR
24984	55352	A	25122	382	1424	VLLENLSEPPNQHDIIEILTVQR
21701	33332	1		302	1 2	KEEETALSCRLVLRKLTETEPVI
						IFEGIEAPATLPADEVPVGKDEN
						DNVEVSRWGTPREFDFEVRDH
						VTLGEMHSGLDFAAAVKLTGS
						RFVVMKGQIARMHRALSQFML
						DLHTEQHGYSENYVPYLVNQD
						TLYGTGQLPKFAGDLFHTRPRH
			1			ECFRHIGSLGLATPLVTECMQW
						LFGIPHTLQLDAIIITCWIILNAIC
						-
						VACGLQKGVRIASDVRSYLSFL
						MLGWVF\IVSGASFIMNYFTDS
		1				VGMLLMYLPRMLFYTDPIAKG
						GFPQGWTVFYWAWWVIYAIQ
		1				MSIFLARISRGRTVRELCFGMV
	<u> </u>	<u> </u>	L	<u>L</u>		QEFGCKSSTKIDNADFNQA

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24985	55353	Α	25123	416	610	
24986	55354	A	25124	662	811	AQQLPGMSDWQNSQEQHQMK N*VRLFMRTSRKLMDRLNVLIK LWSPNRT
24987	55355	В	25125	1	3264	
24988	55356	A	25126	489	3256	PRVFSSATRSLVAMRRWVYRV RRIQYRRTTQTAVFHSTHGNYR EGQPVGTLIDDRYKANENVEL VHYAQPLLNEADSLAKVMPSDI PLKQRRWLGLQMLEGDIYSRA YAGEASQHLDAALARLRNEMD DPALHIADARYQCIAAICDVVS NTLTAEPSRFTTAVDKIVLNRFL GLPIFLFVMYLMFLLAINIGGAL QPLFDVGSVALFVHGIQWIGYT LHFPDWLTIFLAQGLGGGINTV LPLVPQIGMMYLFLF
24989	55357	Α	25127	779	910	
24990	55358	A	25128	2203	2665	ACWARISAAVIQRSGMPGVRV CGRIKSCSKAVRCQRHVFATCV WNGRQVTSGAGTWSTIAWYSG VMSSAISSWSRPAIPFSAEA*TIG KSSC/YVGRVEVNEQIEYLIHNP IRTRARAVNFVDNNNRLQAVS KRFFGYEARLRHRAVKCVNHQ QH
24991	55359	Α	25129	241	918	
24992	55360	Α	25130	2142	2646	KGVCGTTNPAKI\QAILQAFHEI FGEGSCHIASVAVESGVPEQPF GSEETRAGARNRVANARRLLP EADFWVAIEAGIDGDSTFSWVV IENASQRGEARSATLPLPAVILE KVREGEALGPVMSRYTGIDEIG RKEGAIGVFTAGKLTRASVYH QAVILALSPFHNAVY

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24993	55361	A	25131	473	1765	KAWLMTGIVRHWKGRAVPPA EQLLEHLGNGKSFDSVAQAWD AAMADAKAEDTVLWGVEAFRI NAAVAKDDTQIHTHMCYCEFT DIMDSIAALDADVITIETSRSDM ELLESFEEFDYPNEIGPGVYDIH SPNVPSVEWIEALLKKAAKRIP AERLWVNPDCGLKTRGWPETR AALANMVQAAQNLRRGARAG CAADFAASARLAVGEALTNIA ATQIGDIKRIKLSANWMAAAG HPGEDAGLYEAVKAVGEELCP ALGLTIPVGKDSMSMKTRWQE GNEEREMTSPLSLVISAFARVE DVRHTITPQLSTEDNALLLIDLG KGNNALGATALAQVYRQLGD KPADVRDVAQLKGFYDAIQAL VAQRKLLAYHDRSDGGLLVTL AEMAVLSSLGQGGYFGHMII/D
						SGQYPGKSKTGQSTPDVAAQA RQSWIP
24994	55362	A	25132	602	2563	VVTTLTLVLTAVGSVLLLFLV MKARMHAFLALMVVSMGAGL FSGMPLDKIAATMEKGMGGTL GFLAVVVALGAMFGEILHETG AVDQIAVKMLKSFGHSRAHYA IGLAGLVCALPLFFEVAIVLLIS VAFSMARHTGTNLVKLVIPLFA GVAAAAAFLVPGPAPMLLASQ MNADFGWMILIGLCAAIPGMII AGPLWGNFISRYVELHIPDDISE PHLGEGKMPSFGFSLSLILLPLV LVGLKTIAARFVPEGSTAYEWF EFIGHPFTAILVACLVAIYGLAM RQGMPKDKVMEICGHAL\QPA GIILLVIGAGGVFKQVLVDSGV GPALGEALTGMGLPIAITCFVL AAAVRIIQGSATVACLTAVGLV MPVIEQLNYSKRRGFTGRVEEN FGVCKGPSKYHSQHDAINRQG REASDAHGTLENLPLPLVPQPN AATALAALRASGLEVSENAIRD GIASAILPGRFQIVSESPRVIFDV AHNPHAAEYLTGRMKALPKNG RVLAVIGMLHDKDIAGTLAWL KSVVDDWYCAPLEGPRGATAE QLLEHLGNGKSFDSVAQAWDA AMADAKAEDTVLVCGSFHTVA HVMEVIDARRSGGKLCQMHVN VQDKIDEEINQVHICLCLVLVIV SGCKYVGYSLKCSEHYYSLVR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24995	55363	A	25133	518	1079	TQKETNKRCSPLVGWYPAFRM AHFLASERSCYQKLSN\RKSH\P AVAGMVSGMTVANLL/EHSAG NVFKSGI*LALHLFIDRCF*YCG DGIGLFLGARYSRRGERKSART ISLFAQPGPVVNFRRHDVWQRR CVCLVQLRKAIHDVYFRFFGNG DDLYYDVSWARDGAGKYAKW QDFRTLFTTAHCSSD
24996	55364	A	25134	322	798	AQRAHVVGTVGKFDEDDADIL HHRHNHLAKVFGLRLFLIAELE FIKFRHPLYQLGDAFTKKLFHIL ILVVGGSIFNHVMQQRRHQRFI VEFHLGQNAGN\ATGCVI*GSPL ERVCPS*ASRATRYACRRRLIW SIARPAAKVSACCSAIATSK*RS KGQRVLLGNCHVEVAIREFT
24997	55365	A	25135	440	906	CGFSSLSRLPHSSYSAHHAVAPS VLMSASGRPRWSQWYRWIFRA GVYTGSIWFFICSTVRPHDASQ RRLSSLT/QVSQDFTFQPVNHQL WAGGKPTAGITVGKQ*W*PFIK FTVPGDDSGDGRGQGFLQIAAG KCRAQTFFSFCRAGKHDPCRRT VS
24998	55366	A	25136	1059	1541	TAPLLRPGCRLS*SAAHPAAPA QSTNQCPGDRYAATLPAHHRH TNHG*IADVHQHISATPASAAS YAPGFLKDGAPASERHRETGC GRDRPH*PSLYKLSGILRPYQPA SFPG*QQRAVLSAR*LLPVQCSL HSQRCRRRPHRLHRSAPAPGIR ERRSRR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24999	55367	A	25137	236	2211	QGALMSEDIFDAIIVGAGLAGS VAALVLAREGAQVLVIERGNS AGAKNVTGGRLYAHSLEHIIPG FADSAPVERLITHEKLAFMTEK SAMTMDYCNGDETSPSQRSYS VLRSKFDAWLMEQAEEAGAQL ITGIRVDNLVQRDGKVVGVEA DGDVIEAKTVILADGVNSILAE KLGMAKRVKPTDVAVGVKELI
						ELPKSVIEDRFQLQGNQGAACL FAGSPTDGLMGGGFLYTNENT LSLGLVCGLHHLHDAKKSVPQ MLEDFKQHPAV\APLIAGGKLV EYSAHVVPEAGINMLPELVGDL HERSLIFVASGPAGNSKETASD KLEEALPTKNWEVPGANWNTP
						GNGIPSVPEKLKGMLSAPNYRR MVLPAPLQILLLQNAKGGELLG AAGGQIAFNLGSAVGAYCGGM MLTLGLAYNYVALPAALLSFA AMSSLLLYGRYKRQQAADTPV LAKPLGQLDNAPFRCENVNLIR
						EEIGFNALDKFKRTTGALLQLQ QALHPALGADLCGSTGFAVLFI SPVRGNTHLRHLVHIFGTDLDL NRHSVRADHRGVQRLIAVRFW NGDIVFHAARTRLVQAVHLAQ HAITGVRIIDDHAESVDVHDRV KTLLFEHHFAVNRIKMLLPTTD
25000	55368	A	25138		1932	AARYSRFLQTPFDFRKNLLDHL
∠3000	ومحجدا	I ^A	23130	11	1934	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ļ		sequence		
25001	55369	Α	25139	1	1200	MAIPAFGLGTFRLKDDVVISSVI
						TALELGYRAIDTAQIYDNEAAV
						GQAIAESGVPRHELYITTKIWIE
		1				NLSKDKLIPSLKESLQKLRTDY
1		ł				VDLTLIHWPSPNDEVSVEEFMQ
}						ALLEAKKQGLTREIGISNFTIPL
1		l				MEKAIAAVGAENIATNQIELSP
						YLQNRKVVAWAKQHGIHITSY
						MTLAYGKALKDEGIARIAAKH
						NATPAQVILAWAMGGGPVE\W
						RR*QIWRGFSSAS*RHELYITTK
		ŀ				IWIENLSKDKLIPSLKESLQKLR
						TDYVDLTLIHWPSPNDEVSVEE
						FMQALLEAKKQGLTREIGISNF
						TIPLMEKAIAAVGAENIATNQIE
						LSPYLQNRKVVAWAKQHGIHI
						TSYMTLAYGKALKDEGIARIAA
						KHNATPAQVILAWAMGGGPVE
						MAKVADMAGIFVSKLKTFNAG
			}			VKGSTLGASGPQYVMVPKKHA
						DQLRESLASGEIAAVQSTLVAD
						NTPLNSRVYTVRSGDTLSSIASR
						LGVSTKDLQQWNKLRGSKLKP
1						GQSLTIGAGSSAQRLANNSDSIT
						YRVRKGDSLSSIAKRHGVNIKD
ŀ						VMRWNSDTANLQPGDKLTLFV
						KNNNMPDS
25002	55370	A	25140	2237	2404	
25003	55371	Α	25141	307	642	
25004	55372	Α	25142	1449	2084	NAPRCAYY\VDVIAVTMPTTCC
						TLKISPLCRELILTLANRTTTQR
		1				AEPMTRRLIQVLFDELPQQPQQ
						QLHLPVSSHPKIRTMVEMMAK
						GPVEWGALGQWAGFFAMSER
		1				NLARLIVKETGLSFRQWRQQLQ
						LIMALQGLVKGDTVQKVAHTL
						GYDSTTA\FITMLRKPGKTAGTS
						LYHRRYGVSLLSSAWGQVGPP
						RYCRQANSVLSDRFCVLI
25005	55373	Α	25143	981	1103	
25006	55374	Α	25144	222	379	
25007	55375	Α	25145	1	1452	
25008	55376	Α	25146	500	854	
25009	55377	Α	25147	1	1374	
25010	55378	Α	25148	2	167	
25011	55379	Α	25149	781	1047	
25012	55380	Α	25150	1	783	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
25013	55381	A	25151	442	857	LVKNPVKRMEIQQKNLTLFEYL
İ						DWWDVLFLLLESELLFANTWL
		1				AATSESVDDCSGVKPPSGKNGT
						*AMPSWAKASIIASSARCAMLY
						SFCTQTISAILRASATCVGVTLL
						RPICFTRPCCCISASTVRLASIDP
						SAGP
25014	55382	Α	25152	3	4105	
25015	55383	Α	25153	2	269	YKNPYIAAKIQPKTHDTLTPINP
						GRIKLWLMTKRPMWVVPERSN
						*IAARSLG*VGRM**P*QAGAE
						ATMYAGSTPIASATGTIAASAA
		1		1		A
25016	55384	A	25154	1	1370	MCPIEETASSFGGKPLSMVLVIQ
						MFMLLTGALIIILTKTNPASISK
						NEVFRSGMIAIVAVYGIAWMA
						ETMFAPACYGYYILPTYPSDLA
	1					AIQFDRSGTTHIGRFVINHSFILP
						GLIGVSVSCVFGWIFAAIKRDA
						AAGRAKENVIFHHFPFQSVKAD
	}					IATNPFKRPGGAFIRITQTFRTV
		ŀ				QTFRLSAYRLDFAGDRLRISTPR
						AKMRTAFKKDHLRQRRRCIRQ
						RAPPARHNLVGAVALPATVAG
						VNFTFSNVPLDSSVLSSLLTDFS
						TAVGSIVMLAVIMGLMLAFDM
						GGPVNKVAYAFMLICVAQGVY
						TVVAIAAVGICIPPLGMGLATLI
						GRKNFSAEERETGKAALVMGC
		į				V\GVTEGAIPFAAADPLRVIPSI
						MVGSVGGAVTAALVGAQCYA
						GWGGLIVLPVVEGKLGYIAAV
		ì	İ			AVGAVVTAVCVNVLKSLARKN
						GSSTDEKEDDLDLDFEIN
25017	55385	Α	25155	1	789	MKIIAGITPADSGTLEIEGNNYV
					ł	RLTPVHAHQLGIYLVPQEPLLFP
					•	SLSIKENILFGLAKKQLSMQKM
						KNLLAALGCQFDLHSLAGSLD
		1				VADRQMVEILRGLMRDSRILIL
						DEPTASLTPAETERLFSRLQELL
		1	1			ATGVGIVFISHKLPEIRQIADRIS
		1				VMRDGTIALSGKTSELSTDDIIQ
					1	AITPAVREKSLSASQKLWLELP
		1				GNRPQHAAGTPVLTLENLTGE
			1			GFRNVSLTLNAGEILGLAG/TEL
						AETLYGLRTLRGGRIML
25018	55386	A	25156	1	1332	
<u> </u>	1	1	1	1	1	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
			•	sequence		
25019	55387	ĪA	25157	1594	2220	I MWVNPERGRAENTIIGGNATGI
20019				1.52		GTLNVQDQDSVITVRRLS*SAL
						KPGILVFKLTPAETERLFSRLQE
						LLATGVGIVFISHKLPEIRQIAD
					ļ	RISVMRDGTIAGTGFGDDFGFT
						HPLRQQRLPQHLVSFMRAAVQ
						QIFTLEIQRGVRPGCNVLAFGQ
				'		RGWTPGIVFQQVVELSLKFRIFL
			}			RTDKSFFQLAQSRHQDLWHVH
						AAKLTKIĞVK
25020	55388	A	25158	14	237	KTTSFPAKLFNACRISLLASAE/
						CQSEYFSIFSNMDLADTVMFL*
						GAIRFSDLVNEQIQHTFLIPGETI
						CRLSEVV
25021	55389	Α	25159	1	1506	
25022	55390	Α	25160	158	639	SGC*PGRPSLN/GRLDVLSAWS
	i					VSILRLGGMALEGSIAYIVINFC
						DVKPQRFRGLSVPAASITVVLIQ
						QFVLHRRELRRKLALPAHHLVT
						FTWGNVSAVDETRQWMVIKPS
	:					GVEYDVMTADD\RWWLR*PAV
						RWWKAAKNPLPIHQRIWRSTV
						AMPKLAVLC
25023	55391	Α	25161	2	2886	SVPTIIFFFLPVLRFWRPGIQQA
						KKCHLFATGTQSRRNTLRQRT
						AKGIPQQRMTAKREEISIGILHV
						TPQQRRECRGDNRFTARFRRTP
						WRLLGHCVSAAVTGVLPAVAG
		l				SPLAYSDTDEFYPVAGGTMSQ
						HLPLVAAQPGIWMAEKLSELPS
						AWSVAHYVELTGEVDSPLLAR AVVAGLAQADTLRMRFTEDNG
						EVWQWVDDALTFELPEIIDLRT
		ļ				NIDPHGTAQALMQADLPQDLR
					·	VDSGEPLVFHQTTLGFI
25024	55392	A	25162	1	1659	V D S O D I D V I I Q I I E G I I
25025	55393	В	25163	200	646	
25026	55394	Α	25164	1	3197	MPHRQHGPQMTPSHQRTTINR
						DQREQQPIGTAAPQTRPQSAPY
						CEPHTRRRTLQPHNRGFRTAKY
						TSTQARKYHMARPERSNDSDH
						PVHTRSRWNATQTERHTQHHV
		}				SPPSSLRTLRTNCTSHAHHRYH
						RHAQWHRVGLLDVKPTGVPSN
						TVFSPPKNALWADRVDTSRGR
						TTVEQDNHPNGGNVVGTPSDN
,						NFWLETTGGDPNKEYYHHHHY
						NEDYCRYYPLLIAEPLLFPSLSI
						KENILFGLAKKQLSMQKMK
25027	55395	В	25165	1	2505	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25028	55396	A	25166	159	11744	NSRISGLQQRYLMHIMVTQGGL
23028	33390	^	25100	139	11/44	MSLSIVHTRAALGVNAPPITVE
	ŀ					VHISKGLPGLTMVGLPETTVKE
]					ARDRV/RSAIINSGYEYPAKKITI
						NLAPADLPKEGGRYDLPIAIAL
	1					LAASEQLTANKLDEYELVGEL
	1				İ	ALTGALRGVPGAISSATEAIKSG
	1					RKIIVAKDNEDEVGLINGEGCLI
İ	1					ADHLQAVCAFLEGKHALERPK
	1	ĺ				PTDAVSRALQHDLSDVIGQ\EQ
						GKRVLEITAAAGHNLLLIGPPG
			}			TGKTMLASRINGLLPDLSNEEA
ŀ						LESAAILSLVNAESVQKLWRQR
ŀ			•			PFRSPHHSASLTAMVGGGAIPG
						PGEISLAHNGVLFLDELPEFERR
	1	ŀ				TLDALREPIESGQIHLSRTRAKI
						TYPARFQLVAAMNPSPTGHYQ
						GNHNRCTPEQTLRYLNRLSGPF
					ĺ	LDRFDLSLEIPLPPPGILSKTVVP
						GESSATVKQRVMAARERQFKR
						QNKLNAWLDSPEIRQFCKLESE
	1					DAMWLEGTLIHLGLSIRAWOR
						LLKVARTIADIDQSDIITPTYCG
						KADARNSGNVCPSSRALVQR
25029	55397	A	25167	127	503	NRCLLLTLLRCKGRNREEENDA
		`				EAESEKLKNKFNRPSGRLSYGD
	1					RMFDIGVNLTSSQFAKTVMML
						*RARLGLR*TTGLELRELLPLIP
	1					AEKL*/YETDAPYLLPRDLSAKP
						STSRYHCVIEARAWFK
25020	55398	 	25168	3	406	515K11K VIEAKAWEK
25030	122240	Α	23108	J	400	

3331

NO: of p	eptide hod uence	t SEQ ID NO: in USSN 09/540,217	location of first codon for peptide	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
sequ	uence	09/540,217	codon for peptide	lof peptide sequence	Idelation \=nossible nucleotide incertion)
		•			deterion, (-possible nucleotide miser non)
			sequence		
25031 553	399 A	25169	1	1252	MFDIGVNLTSSQFAKDRDDVV
23031	,,,	2310)		1232	ACAFDAGVNGLLITGTNLRESQ
					QAQKLARQYSSCWSTAGVHSH
					DSSQWQAATEEAIIELAAQPEV
					VAIGECGLDFNRNFSTPEEQER
					AFVAQLRIGADLNMPVFMHCR
i					DAHERFMTLLEPQMVENYETLI
					FEAHSTDYQTPQSLRQLVIDHF
		1			AILKVGPALTFALREALFSLAAI
		:			EEELVPAKACSGLRQVLEDVM
					LDRPEYWQSHYHGDGNARRLA
!		ļ			RGYSYSDRVRYYWPDSQIDDA
					FAHLVRNLADSPIPLPLISQYLP
					LQYVKVRSGELQPTPRELIINHI
					QDILAHTTQPVKANKQNKEEH
1					AMPNIVLSRIDERLIHGQVGVQ
					WVGFAGANLVLVANDEVAED
	ļ				PVQQNLMEMVLAEGIAVRFWT
					LQKVIDNIHRAADRQKILLVCK
					T*WKWYWQKGSPYVSGRCKK
	ļ				LSTTFIAPPIDRKSCWFVKHPPIS
25032 554	100 A	25170	2	319	YSSVFSLSASASFSSSRFRPLPT*
	,,,,	25170	[KREKKTPISNRQYIGIANSVCEK
	j				TSGGVSNIPTTNAPTSTYGRFLR
					KSSGEVIPIQHSSTIATGTSKETP
					NAINSAITKLK
25033 554	101 A	25171	1	1011	MIPHPKLEGDRSTQALPPTPVIR
					VTERRNLNWVGTTIGFVHTGK
		ľ			PLSFVYDMADIIKFDTVVPKAF
					EIARRNPGEPDREVRLACRDIFR
					SSKTLAKLIPLIEDVLAAGEIQPP
	İ				APPEDAQPVAIPLPVSLGDAGH
	ļ				RRRRSGRVSEVSWRSGVMSFR
					RASGDKPQPKPKVMPEHAECPP
					AARRGQMMSICPCFALKIDIAD
					NKFFNGETSPLFSQSQAKLARQ
					FHQKIAGYRPTPLCALDDLANL
		1			FGVKKILVKDESKRFGLNAFK
					MLGAPEKCHRRKNDF*RRSDRI
					VLIRL*REFSSHQKGLRDRPAH
					KAVLGDNQLFSGGTGAPVWPG
					FAKRAVMF/TPLKNLLSAISIFN
					AKHGHILIIWPRLAAGGHSACS
					GITFGFGCGLSPLARRKLMTPE
					RQLTSDTRPDRLLRWPASPSET
					GSGMATGCASSGGAGGCISPA
				· '	ASTSSISGINLANVLLLRKISLH
]]					AKRTSRSGSPGLRRAISKAFGTT
		1			VSNLMMSAIS
	1	ŀ		l .	
25034 554	102 A	25172	1	510	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25036	55404	A	25174	704	1069	RLWQADCKENAG\GDYYWTFF PQPTFVSTQKYYCHVDNSCYM NFDFSAPEYHAHWRYGLLMVG ESQHGLGANAIQASQCFKAAN QPLGADATFLQQIKGEIICFGLFI TGVFSIDSAFCDIT
25037	55405	A	25175	1	639	MARVWLCDFIPYGYLPTLNAM PESTRKQIMPWTRLSGSLHTWN RTPNRRIGRYPSGLPPDSANRPR REPDACRSLHSPAKYRGNPNRD RLDHQQNGLPARGKDTPRSNK PSISQWTNTKPQPRHQPRGKKG HKSPAPEMPKPPQATSSTREQT KYGAAELWRSPWKTRYIELKA RGPMACVTISLNVCCAHLVRL VMVTLKQTDPIRNLGEMPSKSQ TKKAREVHLHSLHLQELAQEQ EPPARKRKTQWAVSERK
25038	55406	A	25176	82	356	AQCEEYRVPQTLKRNPKQCFDI SIRGGSQPCGIFSTPVRNTLQNM GQMGWLVAPG*WLWREIRKH NRWCACPRKPGRAGISMALRW GSGHA

NO: of peptide sequence hod in USSN location of first codon for last amino acid of peptide sequence sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25039 55407 A 25177 1 2635	MNKSGKYLVWTVLSIMGAFAL GYIALNRGEQINALWIVVASVC IYLIAYRFYGLYIAKNVLAVDP TRMTPAVRHNDGLDYVPTDKK VLFGHHFAAIAGAGPLVGPVLA AQMGYLPGMIWLLAGVVLAG AVQDFMVLFVSTRRDGRSLGE LVKEEMGPTAGVIALVACFMI MVIILAVLAMIVVKALTHSPWG TYTVAFTIPLALFMGIYLRYLRP GRIGEVSVIGLVFLIFAIISGGWV AESPTWAPYFDFTGVQLTWML RLINRSIMMDFSLTEEQELLLAS IRELITTNFPEEYFRTCDQNGTY PREFMRALADNGISMLGVPEEF GGIPADYVTQMLALMEVSKCG APAFLITNGQCIHSMRRFGSAE QLRKTAESTLETGDPAYALALT EPGAGSDNNSATTTYTRKNGK VYINGQKTFITGAKEYPYMLVL ARDPQPKDPKKAFTLWWVDSS KPGIKINPLHKIGWHMLSTCEV YLDNV*TSVWRTCACPALRRA TQRYRRQSRAGNLRTSGEAIRR DMEWTSLRSE\DWEKGDTIKQC ISAATSCVYGVTGEGILEGGDA QILREPLPGTKLGLLNWPVFDA LESRSTRERDGKPGGEICGETA NILIRGPENRPLRRDPPGKP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	<u> </u>
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25040	55408	A	25178	216	1829	TVCRCHFVTTKHIGRERNQRAR
						QCLVGKTTDTQQHNCDIRVGE
						QADHTRANHQHCTQRYDHLTG
		1				NDQVHTKPTLEQRRQVTTEDT
						TEVSKQHRHPGEHRDLFQRTLN
İ					·	ISADDORRLLLELQNDNLNHNR
		ļ				IWLRLAAQPEDHIYGCGEQFSY
						FDLRGKPFPLWTSEQGVGRNK
						QTYVTWQADCKENAGGDYYW
			}			TFFPQPTFVSTQKYYCHVDNSC
		İ				YMNFDFSAPEYHELALWEDKA
						TLRFECADTYISLLEKLTALLGR
						QPELPDWIYDGVTLGIQGGTEV
						CQKKLDTMRNAGVKVNGIWA
						QDWSGIRMTSFGKRVMWNWK
						WNSENYPQLDSRIKQWNQEGV
						QFLAYINPYVASDKDLCEEAAQ
		İ				HGYLAKDASGGDYLVEFGEFY
						GGVVDLTNPEAYAWFKEVIKK
						NMIELGCGGWMADFGEYLPTD
		1				TYLHNGVSAEIMHNAWPALW
						AKCNYEALEETGKLGEILFFMR
						AGST/GDIGG\YTTLFEMKRSKE
			1			LVLRWCDFSAFTPMMRTHEGN
						RPGEHWAF/DGDAKHRHLAVY
25041	55409	A	25179	277	488	
25042	55410	В	25180	260	1500	
25043	55411	Α	25181	1	764	MQEWGRYTILSGCCNSALKGD
						VCLLYGFRVAGRRQASE*TPGA
						YNRVPSSKK*GFQLVLLPFSAN
						GKSPAY*RFSKSFAEDGEKDAG
		1				FPHPYAEAGYDNGGRRSGAGT
						LLSR*I*TGAGRGAWRV*RLTG
						E*MAGKRT*VTASGYSQR*IMV
						FLWSLYRSYRL/GRVRQHNGPR
						YLPVLARNWKMSAWVAAAW
						QGLTDMKRKTMKIRSGS\IVILA
						SGYAATAA*PLSGDRIFLP*PGK
						TG*RHGGTPSCAGFTGDY
25044	55412	В	25182	1	3117	
25045	55413	В	25183	270	388	
25046	55414	A	25184	2154	2279	DDNGC AD # CD CD VD CT OC VD CT
25047	55415	Α	25185	614	623	RPNSGAR*GRGDKRSTQSVPET
		1				VRLLFCAGTFDQQPGNARWDD
						QYRCIRSGIVGVRAVLLGGDIL
L	<u> </u>	<u>l</u>		<u> </u>		DTQPLPVELAETLGKSAE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25048	55416	A	25186	1208	2813	CYGVAWPFDPTLKFEFWRYFT
1						HALMHFSLMHILFNLLWWWY
1						LGGAVEKRLGSGKLIVITLISAL
}		1				LSGYVQQKFSGPWFGGLSGVV
		ŀ				YALMGYVWLRGERDPQSGIYL
				ļ		RRGVIIFAPIWIVRRWFEMTPTQ
						WSNYFFENLFKYEWVQTRSPA
	1	ŀ				GAIQFEAVDAPEIIPDPFDPSKK
						RKPTMLVTDLTLRFDPEFEKISR
						RFLNDPQAFNEAFARAWFKLT
						HRDMGPKSRYIGPEVPKEDLIW
						QDPLPQPIYNPTEQDII\DLKFAI
						ADSGLSVSELVSVAWASASTFR
						GGDKRGGANGARLAL\MPQTR
						TGDVTRRQPIRAL\PVLE\KIQKE
						SGKASLADIIVLP\GLFVVEKSA
						SAAGLSIHVPFAPGRVDARQDQ
						\TAIGMFELLEPIADGFRNYRAR
						LDVSTTESLLIDKAQQLTLTAPE
•						MTALVGGMRVLGANFDGSKN
						GVFTDRVGVLSNDFFVNLLDM
						RYEWKATDESKELFEGRDRET
						GEVKFTASRADLVFGSNSVLRA
					ł	VAEVYASSDAHEKFVKDFVAA
						WVKVMNLDRFDLL
25049	55417	A	25187	1520	1960	PVPLNFGWQRSTEVLICHVNSP
						STAAMTISPWRR**RRRTKQLS
	1					APLLLGVSAIGW/LTIILVAFMA
						WLLAKT/AFGRSFYATGDNLQG
						AR/QLGVRTEAIRIVAFSLNAER
		1				DFDDLRDSGPLIAMILFIMPTLS
						SISSRRLNPGVPSEI
25050	55418	Α	25188	1155	1778	SASGGVMLKFIQNNREITALLA
		1				VVLLFVLPGFLDRQYLSVQTLT
						MVYSSAQILILLAMGATLVMLT
						RNIDVSVGSITGMCAVLLGMLL
						NAGYSLPVACVATLLLGLLAGF
						FNGVLVAWLK/IPAIVATLGT\L
						YRGLMSLWTGGHWFEGYPRIE
}		Ì				KPSPPVAGVHHWCVTQIGGLRP
						AVAKRVGPDLPGDNKGAKGVL
			1			NTLDNYGVDGKCV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25051	55419	A	25189	1	2227	I MVTIAGIILGGLALVGLITYFGK
25051	33419		2310)			WTYLWKEWLTSVDHKRLGIM
						YIIVAIVMLLRGFADAIMMRSQ
						QALASAGEAGFLPPHHYDQIFT
ļ						AHGVIMIFFVAMPFVIGLMNLV
						VPLQIGARDVAFPFLNNLSFWF
						TVVGVILVNVSLGVGEFAQTG
						WLAYPPLSGIEYSPGVGVDYWI
						WSLQLSGIAVAHIAAQNVAVLL
						ISSDLEEIELMADRVYVMHQGE
						ITHSALTERDINVETIMRVAFGD
						SQRGVMLKFIQNNREITALLAV
						VLLFVLPGFLDRQYLSVQTLTM
						VYSSAQILILLAMGATLVMLTR
						NIDVSVGSITGMCAVLLGMLLN
						AGYSLPVACVATLLLGLLAGFF
						NGVLVAWLKIPAIV\ATLGTLG\
						LYRSIMLLWTGGQWIEGLPAEL
						KQLSAPLLLGVSAIGWLTIILVA
						FMAWLLAKTAFGRSFYATGDN
						LQGARQLGVRTEAIRIVAFSLN
İ		İ				GCMAALAGIVFASQIGFIPNQT
						GTGLEMKAIAACVLGGISLLGG
						SGAIIGAVLGAWFLTQIDSVLV
						LLRIPAWWNDFIAGLVLLAVLV
						FDGRLRCALERNLRRQKYARF
						MTPPPSVKPASSATSDFICIGIVA
		l				LPLTMVIVSGGIDISFGSTIGLCA
						IALGVLFQSGVPMPLAILLTLLL
						GALCGLINAGLIIYTKVNPLVIT
						LGTLYLFAGSALMLSEEQKNTK
						KTRQREDKKKEKKKQRKREEK
25052	55420	Α	25190	1055	1206	SFALFSSSSPRSRP*FASTKS*TL
						TTGQVLPSTNVRFLTGNWGPPV
		<u>L</u>				VATG
25053	55421	В	25191	1	1386	
25054	55422	Α	25192	1	449	MQDPLLALLEDYNKGDLEKQV
						PPAQSGPQPGKAGRDRPIVIML
						DPGHGGEDSGAVGKYKTREKD
		l				VVLQIARRLRSLIEKEGNMKVY
						MTRNEDIFIPLQVRV/AKAQKQ
		1				RADLFVSIHADAFTSRQPSGSSV
		ļ	05100	1506	1041	FALSTKGGMCSYRGSGCLQ
25055	55423	Α	25193	1536	1941	STSNQNSTALCRRRGLVHCIDG
						ANDDDSVMVWPSGSGRPWSAE
						PGAPQR\YAQLVVEDVELDLLH
						FIVFGDDVMLNVVLQGQAFML
						ALGLVTNVGVLAHAHHHALVS
1						GVPNDGGGTWVGGRCVQQSQ
L	<u></u>	<u> </u>		<u> </u>	<u> </u>	LCTCWSHCQ

NO:				Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
1,.0.	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25056	55424	A	25194	240	855	SRAGYAPAAAQKVASPYVRRE
						R*NQSD\PAVLTGIGERGK*LLS
						RDFFIAPLAGMIVEKRGAVHLA
			}			RWSVPVERKGQWQPAGLWAQ
						FFLADIVCPAATALPDTTAEDQ
						HIDHPAVVHVHVIPVVNTCAED
						NHRAPMRFMGGIGKFTGDGFNI
		l				LARHPGNLLTPGWGVRFDFSVI
						FRTVFIFQPAIEAVIRQYQIINAD
						HRTGVAID
25057	55425	A	25195	840	1157	QYAPTAQPARPADPFALPTVQA
2000,	55 125	ļ · ·				RLTPFCYLVAKGDWCVYRRSY
						SLWHSVASPRVRQSPRLGY*GY
						TANSPGILLAPLSLRGEMFLPOD
		ŀ				APKSVPPYKRGKSVAIN
25058	55426	A	25196	1076	1339	LVCVASMPVEISTDASASSGVI
23030	33 120	l'`	23170	10,0		VISSAMSGSWSAETKRRLAMV*
						PEPLHLWAISIPTTSGRFTSSLIS
						PFPAR*YSPARYIPRVWRV
25059	55427	A	25197	2	118	THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL CONTROL OF THE TOTAL
	55428	Α	25198	1809	3145	ARHKQHAYGYRYRQSPRCYGV
25000	33 120	1	25170	1005		SGKKRGLRYYGSWLAISGSSRI
						TR/LLCMYLRRFFIGVIGADIAN
						MGISAGAMWLLSVSQVSLAAV
		l				SQVVAVRVWPASSYTRVTVES
						NRQLKYKQFALSNPERVVVDIE
						DVNLNSVLKGMAAQIRADDPFI
						KSARVGQFDPQTVRMVFELKP
						NVKPQLFALAPVAGFKERLVM
						DLYPANAQDMQDPLLALLEDY
						NKGDLEKQVPPAQSGPQPGKA
						GRDRPIVIMLDPGHGGEDSGAV
		l				GKYKTREKDVVLQIARRLRSLI
		l				EKEGNMKVYMTRNEDIFIPLQV
		l				RVAKAQKQRADLFVSIHADAF
		l				TSRQPSGSSVFALSIQRGARPVG
						DRYVDHTMFDMVQSLTIADSL
						KFGKAVLNKLGKINKLHKNQV
						EQAGFAVLKAPDIPSILVETAFI
						SNVEEERPNGQPGHPLDPATTM
						GTLIDCAHADSVQ
25061	55429	A	25199	1	277	G I BIDCAHAD3 VQ
	55430	A	25200	902	1006	
	55431	A	25201	2	1265	
	55432	A	25202	409	525	
	55433	A	25203	91	431	NHKPGNIDVARRIQRGFAGDQI
			•			GHLRPVERQCSPDKRRFIAADG
						REIRGKQRAGHIFQLLSRCLLQI
						LNHCQRRAAHFRFQLSNQRHQ
						QLLPV/HYHAAEREYPAGACLV
						RWLL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25066	55434	Α	25204	3	71	
25067	55435	A	25205	1	1212	
25068	55436	A	25206	1	2786	MAAVIEQIRRAVLALVTGVVHP GDIPVVWDIVWNATAAFIAVIII SLLLDESGFFEWAALHVLRWG YGPGRLPFNRIGLIRRITRRIRQR LLFGLVTFILLLNRSQNPAQDIQ LLAFRARTGKQTAQFIHHLPRM VFTDKTGSSNGLAPQGQCSAQ GELILNEKLAKQLVTAANWVK MQSDEGEINPVDILRWPGVMA AQEQDLDAIAAEILAALDGTLD DFIVARETEGQSGLKRVYHSPG
25069	55437	A	25207	5	402	APDIREFTRDAIP EIFSVVWIIMTRGDVVTISRWR VSSSPLETWNKRWAKISPAPFS LLPISRLEKSEGDWLPETVISAF NMPSSLSLIISSL*FKVRDV*HFF /RI*TLRGHYRVINWPNFNIVVS QGIGKRLANSWLVFTLFHRI
25070	55438	A	25208	48	219	PETCGHLWAYVWPSCAAV\GL YFKVHVLG*RSVTPVTDIVKLL EFTRLRLPGYTKSIE
25071	55439	A	25209	1	912	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25072	55440	A	25210	<u>I</u>	1857	MRVCARACVRTRTRMCVYAH
23012	33440	^	23210	1	1657	TCVCVRTYAYVRVRVRVHVRA
						RVRVCAYARTRVRNSLSILPFIQ
						LTLATPIHHIHQEEFNIRGIVPVL
						RRVKPDLAIGIDITPSCDTPDLH
						DYSEVRINQGVGITCLNYHGRG
						TLAGLITPPRLIRMLEQTALEHN
						IPVQREVAPGVITETGYIQLFLP
						GWEIGFSPLALLLAFLCSTSPGF
						GDPDGLGVIAYQDTVRPNAAT
						AISELNALAVKGVILTGDNPRA
						AAAIAGELGLEFKAGLLPEDKV
						KAVTELNQHAPLAMVGDGIND
						APAMKAAAIGIAMGSGTDVAL
						ETADAALTHNHLRGLVQMIEL
						ARATHANIRQNITIALGLKGIFL
						VTTLDDRVVAGSAGRYGGDGA
						GDSECVKIVAQEIRQTDRNIER
						DRSPRPSGERVRVRGKGGIEAN
						QPLSTAFTNQITIRQSIRLFSNQF
						VFHKGISRVVAGSSAIISNVSPG
					}	CILRTPNCRRGFPISCTGEGKAA
					Ì	YGWQRDGGHAEYLLAEEKDLI
						LLPDALSYEDGAFISCGVGTAY
	1					EGILRGEVSGSDNVLVVGLGPV
						GMMAMMLAKGRGAKRIIGVD
						MLPERLAMAKQLGVMDHGY*I
1						LPDVY/V*IGVARVSWMNGRID
						SELRTRVRAYAHTRTRARTCTR
			ľ			TRTRTYAYVRTHTHVCAYTHM
						RVRVRTHARAHTR
25073	55441	A	25211	3	1180	SKLGTRRSVVWA*SPSTSPTLW
23073	33	ĺ`.				CSTFSAAGHSSMKRMNEFVDL
		ł				LPAQQRMKGENWYRGTADAV
						TQNLDIIRRYKAEYVVILAGDHI
		İ				YKQDYSRMLIDHVEKGARCTV
		1				ACMPVPIEEASAFGVMAVDEN
						DKIIEFVEKPANPPSMPNDPSKS
						LASMGIYVFDADYLYELLEEDD
						RDENSSHDFGKDLIPKITEAGLA
		ļ				YAHPFPLSCVQSDPDAEPYLAR
						CGYAGNFMESEPRSGLIRVSCM
		1				PAGPPLPLPAARSEAPKAAGTV
				1		ASVPSIAPARLRPVPDPVELVLV
		1				AAEFITPGDPTPRLHGSGFIDIR
		ĺ				QIHHQTRSHLEGVKTGIRFLNH
						FSGNPQGGIAHVNGVARFQVK
						QCHQAWGQQYAARLRFQARGI
						SLQIAIHRVDIIHRFDVRQL
25074	55442	Α	25212	16	543	
25075	55443	Α	25213	1	3387	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25076	55444	A	25214	11688	2234	LSVGKTFPVLVPAAFRKVSATG
25070				1.000		AAVPGTPSILALPSYSTTPFGVC
						TSASMNVPVLTLRIRTRVTGEV
			i			AAIKPFSMANAPTPDSILPQLGV
						VSTRCS*TITCANR*STSARGSL
		Ì				ERLIMATLL/WVAGTFD*SVWF
						GHDTDVVTLYRFHEALCYSVT
						LRTPHRRVLRFKSQHPGELACFI
						SPVA
25077	55445	A	25215	905	4761	PPHNWMPSNATPGIAFVWCAY
						GAI/LPGDAPVPVVDDYRKVVR
						KDTKGLIARWKYFWMSVIALG
			.			VAFALYLAGKDTPATQLVVPFF
İ						KDVMPQLGLFYILLAYFVIVGT
						GNAVNLTDGLDGLAIMPTVFV
						AGGFALVAWATGNMNFASYL
						HIPYLRHAGELVIVCTAIVGAG
						LGFLWFNTYPAQVFMGDVGSL
		E				ALGGALGIIAVLLRQEFLLVIM
		ŀ				GGVFVVETLSVILQVGSFKLRG
						QRIFRMAPIHHHYELKGWP
25078	55446	Α	25216	269	1219	QTRTRRVQLQDLIEYVAIHQW/
						FSDKHPVPFYSCIFDDSITNCYS
						WSISKDGYFIFGGAYPMKDGQT
						RFLTMKDKMSAFQFQFGKTVK
					ł	SEKCTVLFPSRGQDFVCGKDNA
		:				FLIGEAAGFISASSLEGIIYALYS
						TDIPRSVLLKQPDKLNPAYWRA
					:	TRLVCITLFGQIVNTICMAAPDL
		İ				RKWIMLSPVAHKGHHLEWVLH
		1				DMQILPVPRPSGCICSGSSTFFC
						NWGCPYDRPQWQLYPSMCLC
						VHIGPAPVRSEKVESKRAISKV
		l				DQD\$AFRGCCRSGNRNGGRKI
						VAVSLKKVSGAAKPCGHFRST
						RYRVAYSTIIN
25079	55447	Α	25217	l .	1158	
25080	55448	Α	25218	1156	1899	LKLGIRYPQCQRLVTHSLDGISL
						PLHTAH*SQSNSAENTHRAISLK
						AVPPFFTAERRGVNAEEHRNGR
						LINGECRQRFNVLRVANGVRN
						VQFAKARDRNDIASFRQIGFDT
						FQTEVAQHFTDFRVAGFAFAID
						DSDLLVRLHFTALDAANADNA
						NIAVVIELRNLHLERTVKVNVR
						RRNVVNNRLVQRGHVFRHIFV
						VQTRDTVQRRSVNDREVQLLV
						GRVEVNEQIEYYADAFVIMLER
		<u> </u>				LCKVLCCR
25081	55449	Α	25219	1	2583	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25082	55450	A	25220	278	787	TLLSATRQVYALLSTKPAFFSS HSSFSTSSRLISSRAASSMDVAF PIAFSASKTVSIKFFSR*KLLRNS HRPPFMVMRSIIRSPSNVFMAS* FRFVSIPTLRIDAVRRSTIWCSTT ARFSSRADAKFIKNVFQQAIPDS AAKWFFPVPGFPLNAKFVHPVI SSSAMDFA
25083	55451	A	25221	866	1144	
25084	55452	A	25222	19	1203	NWNLTPYASLTGFTADNPEYH LSNGMKSKSVDTRSIYRELGAT LSYNMRLGNGMEVEPWLKAA VRKEFVDDNRVKVNSDGNFVN YLSGRRGIYQAGKDLDRFKNL VLVHAARYAADLSYLPLMQEL EKRYEGKLRIQTVVSRETAAGS LTGRIPALIESGELESTIGLPMN KETSHVMLCGNPQMTRRLIPAS VEVYHDSLC\RKIWHGWEQHIT ARYLVGADGANSMVRRHLYP DHQIRKYVAIQQWFAEKHPVPF YSCIFDNSITNCYSWSISKDGYF IFCGAYPMKDGQTRFTTLKEK MSAFQFQFGKTVKSEKCTVLFP SRWQDFVCGKDNAFLIGEAAG FISASSLEGISYALDSTDILRSAL RKPIRTEQGANGAIRELEKFRPA
25085	55453	В	25223	58	359	
25086	55454	Α	25224	4	329	
25087	55455	Α	25225	1	1008	
25088	55456	A	25226	1	1335	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25089	55457	A	25227	3	1886	HRLNITAENDCRRLHCSLRDLS
						SLLQAVGRLAEYFIGDVFAARF
						NDALTVVERLVKVTLYGSQIKL
		İ				YNIETAVPSVLKPDLIDVHAQS
						LAGGCRLSAHWLAQYCSEIHR
						QNTQQFVTLISTTMDAITPLIST
						KVQDKLLLSACHLLVSLATTVR
						PVFLISIPAVQKVFNRITDASAL
						RLVDKAQVLVCRALSNILLLPW
						PNLPENEQQWPVRSINHASLISA
						LSRDYRNLK\PSAGAPQRKMPL
						DDTKLIIHQTLQRLRKILWENIS
						G\ESTKSSTDFAYQFGCRESV\Q
						VSLGLFPAF\IHQSDVTDEMLSF
						FLTLFRGLRVQMGVPFTEQIIQT
						FLNMFTREQLAESILHEGSTGC
						RVVEKFLKILQVVVQEPGQVFK
						PFLPSIIALCMEQVYPIIAERPSP
						DVKAELFELLFRTLHHNWRYFF
						KSTVLASVQRGIA\EEQMENEP
						QFSAIMQAFGQSFLQPDIHLFK
	ŀ					QNLF\YL\ETLNTKQKL\YHKK\I
						FR\TAM\LFQFVNVLLQ\VL\VH
						KSHDLL\QE\EIGHRHSYNMASV
						DF\DGFFAAFLPEFL\TSC\DGV\
						DANQKSVLGRNF\KMDRLCPCL
1						PGPAPHSPRMCTGWSTTCGLLT
						DSGNDSLAPWGLWKLLGPCLL
}						PGGTRDFLLVAT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25090	55458	Α	25228	689	3053	VDCRMS/FSYIEKFTDFLRLFVS
		•				VHLRRIESYSQFPVVEFLTLLFK
		l				YTFHQPTHEGYFSCLDIWTLFL
}						DYLTSKIKSRLGDKEAVLNRYE
						DALVLLLTEVLNRIQFRYNQAQ
						\LEEL\DDETLDDDQQTEWQRY
						LRHSLEVVAKVMELLPTHAFST
						LFPVLQDNL\EVYLGLQQFIVTS
	,			}		GSGHRLNITAENDCRRLHCSLR
						DLSSLLQAVGRLAEYFIGDVFA
						AR\FNDALTVVERLVKVTLYGS
1						QIKLYNIETAVPSVLKPDLIDVH
						AQSLAALQAYSHWLAQYCSEV
						HRQNTQQFVTLISTTMDAITPLI
						STKVQDKLLLSACHLLVSLATT
						VRPVFLISIPAVQKVFNRITDAS
						ALRLVDKAQVLVCRALSNILLL
						PWPNLPENEQQWPVRSINHASL
						ISALSRDYRNLKPSAVAPQRKM
						PLDDTKLIIHQTLSVLEDIVENIS
						GESTKSRQICYQSLQESVQVSL
						ALFPAFIHQSDVTDEMLSFFLTL
						FRGLRVQMGVPFTEQIIQTFLN
			ŀ			MFTREQLAESILHEGSTGCRVV
						EKFLKILQVVVQEPGQVFKPFL
						PSIIALCMEQVYPIIAERPSPDVK
						AELFELLFRTLHHNWRYFFKST
						VLASVQRGIAEEQMENEPQFSA
						IMQAFGQSFLQPDIHLFKQNLF
		I				YLETLNTKQKLYHKKIFRTAML
1						FQFVNVLLQVLVHKSHDLLQE
						EIGIAIYNMASVDFDGFFAAFLP
25091	55459	Α	25229	1	853	MISVANEIAEAGFDPQGRTSED
						LLDLAESRVFKIAESRANKDEG
						PKNIADVLDATVARIEQLFQQP
						HDGVTGVNTGYDDLNKKTAG
						LQPSDLIIVAARPSMGKTTFAM
						TLVENAGR\MLQDKP\VLIFSLE
						MPSEQIMMRSLASLSRVDQTKI
						RTGQLDDEDWARISGTMGILLE
						KRNIYIDDSSGLTPTEVRSRARR
ļ		1				I\AREHGGIGLIMIDYLPLMRVP
						ALFDNRTLEIAEISRSLKALAKE
						LNVPVVALSHLNRSLEQRAEER
						PGNSDLGESGSIEQDADLIL
<u> </u>			1	<u> </u>	L	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25092	55460	A	25230	206	1073	VKITSWLMSNSALPGTLKLLAL
23092	33400	^	23230	200	1075	ILKCSTCVRVLPRSELRNLER/Q
						GQLDSVGGFA/YLAEL/SKNTPS
		1				AANISA/YADIVRERAVVREMIS
1					,	V/ANEIAEAGFDPQGRTS/EDLL
		1				DLAESRVFKIAES/RANKDEGR
						NPMFDDRGKRGEPSDLIIVAAR
						PS/MGKTTFAMNLVENAAML/Q
		1				DKPVLIFSQTKI/RTGQLDDEDW
						ARISGT/MGILLEKRNIYIDDSSG
						L/TPTEVRSRARRIAREH/GGIGL
		l				!
						IMIDYLQLMRVP/ALSDNRTLEI
	ŀ					AEISRSLKHFEKLALICGCRNNG
			2 - 2 2 1		100	TTRYCKSLLA
25093	55461	В	25231	287	483 635	
25094	55462	В	25232	· · · · · · · · · · · · · · · · · · ·	2469	MPSDISLASCRLRATSRLFTDHR
25095	55463	A	25233	1	2409	RQSDCMTAINRIPIVDDEDNVR
						RETHCANNVRTALHLEADIHP
i						DVVLMDIRMPEMDGIKALKEM
						RSHETRTPVILMTAYAEVETAV
						EALRCGAFDYVIKPFDLDELNLI
]		l				
					;	VQRALQLQSMKKEIRHLHQAL
·						STSWQWGHILTNSPAMMDICK
						DTAKIALSQASVLISGESGTGKE
						LIARAIHYNSRRAKGAFIKVNC
						AALRESLLESELFGHEKGLFER
1	İ					ANEGTLLLDEIGRM/PLVLQAIL
						RILQEREFERIG/GHQTIK/VDIAS
						LLAPTRLQAMVKEGTFREDLFY
						RLNVIHLILPPLRDRREDISLLA
						NHFLQKFSSENQRDIIDIDPMA
	}					MSLLTA WSWPGNIRELSNVIER
						AVVMNSGPIIFSEDLPPQIRQPV CNAGEVKTAPVGERNLKEEIKR
						VEKRIIMEVLEQQEGNRTRTAL
						MLGISRRALMYKLQEYDATGF
						FRDGMTIMVGGFMGIGTPSRLV
						EALLESGVRDLTLIANDTAFVD
						TGIGPLIVNGRVRKVIASHIGTN
						PETGRRMISGEMDVVLVPQGTL
						IEQIRCGGAGLGGFLNPTGVGT
						VVEERQTSIDTSTRKRPFELGRP
						SRNNDKGIRPRSGAGSEEYPRT
						LHDSLKRLQTDYLDLYQVHWP
		-				LRPFYCFGKLGYSWTDSAPAVS
						QLDTLDALAEYQRAGKIRYIGV
						SNETAFGVMRYLHLADKHDLP
25096	55464	Α	25234	1	891	
25097	55465	Α	25235	1	1614	
25098	55466	Α	25236	1	946	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25099	55467	Α	25237	1055	1474	NGVKSKPPPNHQTGSAPGFSAI
						KNRTFAWPTLPNTSCWANTML
		ļ				IGSRELGIRSSQSGPLLGVRVWC
	1	l				/CRCRGTQG/HSQGKDG/IPENGI
						RAYKFTSS*ASFRNSACFRQPSR
						IKPLRKATWLLCGLSTRHLITK
						RLSSPDWA
25100	55468	Α	25238	2036	2743	DLLIRQMPLLSRRVIPVIRVPVT
						GVLWKSFTPRITIFGH/ALIDQH
						SMIVAAQGLTPDHQLLLQIYDR
	1					ARVSASRIVHQAQIYGDAVVR
						YAFIEHRAEVFDFASIEGNEENN
			ļ			VWLCDCAKVYGHAQVKAGIEE
						DAIPTIHYSSQVAEYAIVEGNCV
ŀ						LKHHVLVGGNAVVRGGPILLD
						EHVVIQGESRITGAVIIENHVEL
						TDHAVVEAFDGDTVHVRGPKV
						INGEERITRTPLAGLL
25101	55469	Α	25239	2234	2571	IMSSVDSSLVLPLKAMVPSRITL
						IRSAIWRISG/RLMRDKNNTHAS
		l				SKQLLQATKQAFGFRRGKRGG
		ł			[RFIEDQNPRVAHQPAEDFHHLA
		Ì				IGDIQRSCQTMQIKLAAQRRQQ
	<u> </u>	ļ				VLH
25102	55470	Α	25240	659	1206	WKSTSNTAKAKPGSVMAVDLR
						IPIRFGVMLSAA/FGDDVVDLIA
						QVDLVTTAVGPVVLERIAPAIA
					-	KGQVKRKEQGHESPLNIIACEN
						MVRGTTQLKGHVMNALPEDA
						KAWVEEHVGFVDSAVDRIVPP
						SASATNDPLEVTVETFSEWIVD
						KTQFKGALPNIPGMELTDNLM AFVERKLFTLKH
25103	55471	A	25241	332	683	ALPGR*RHVHLPGSAYEWNDA
23103	334/1	A	23241	332	083	EPAQESDELSLKMLTAICWLTA
						TA/STSLKI*R*KVALRCLVGRIR
						RLRRIRQQCQLLMRRSRVLSGR
						LSSV
25104	55472	A	25242	1	841	MVKVAGECRVRASAHLPGLNR
23104	33472	^	23242	1	041	STKNTEDRGANNQEGTNTQKT
1		l				RGETNERLKKNEHRDLRLARY
1						PAKENNKQGDKNKPRTYEKLT
						GTVTHQEPPHHADQSPGLCLFR
						TSRAQVERGPTVLLLISQENQIE
						RYDSDNHSQLNLCLRVPTVTTQ
				1		RRDYDIAERYRQKGIKLPARKE
						RLMTIAERLRQEGHQIGWQEG
						KLEGLHEQAIKIALRMLEQGFD
						RDQGIA/RRASCQRQQRQSSLPT
						P/EHCCAISGATARKPAALPAAR
						IRRILVSIIRCTSGCPALSG
L	<u> </u>	<u> </u>	<u> </u>	<u> </u>	l	INGL VOINCEDUCE ALSO

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
1		ł		sequence		
25105	55473	Α	25243	240	651	RCPPQSCLLAEGRWCG*K/CRK
	1	1				VSV*SNRYR*SRLSLGELLRAYS
		ł				RNRDLPGF/SSGSARVLVATDV
						AARGLDIKSLELVVNFELAWEP
						EVNVHRIGRTARAGNSGLAISF
						CAPEEAQRANITSRPPHPAHIAS
						DHQSTP
25106	55474	Α	25244	1455	1643	RDPHQICVAESVYLAAFVLLCA
						GLCVRRRSTTGARNAAAAC*N
						GLW*AEAQCPAYYVCPVVG
25107	55475	Α	25245	549	641	
25108	55476	Α	25246	1378	2084	PNFGYSPPASSCGQVSQNSS*LQ
İ						CSVSGRDKMLRTGPCARTVSSS
						YLMGVSFSRDA**DAR*VCACH
						NTIAALSASAERFTSCERASHSE
						ARSGCCNSVRVMSNNPCCPRSL
						IGVYDILFFQESTAKRCKRSLSG
						SGSACGKSRVRFTPYRPARRTP
						LRSTTSPISAASFPSGWECNAAS
		l				TCSAALRSTAITSLPSLAKFRAS
						SPSNSQTPRTAGLTGSAASESSI
						PQPQAAANS
25109	55477	Α	25247	1134	1428	GPCGCCSSNVARCASQTSSRFA
						AV/WHGAIGAR*LNADNRPLWI
						EITREHAELPGKTGRRMETKQR
						RAITFTQRQNGFESHTPGAVFFS
						INNRCQFGDR
25110	55478	Α	25248	96	535	RPQTTVQPALYVAEQSICSQPC*
						*FSTKCDRKPAAVLAPAGYAAL
		ł				PVLFSSAKLDGSWLRWNSASG
						RRQNGSFSCAALASKRPASASS
		İ				KLNSA**SLPSAVFAAPVRVAA
						SMIISGFCALASIRPSASTRRPSA
						SVFITSTFLPLR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25111	55479	A	25249	1471	3026	SPLRSSVACRGWHRKNNRRGW CPRRGDPARPVRDSLQ\HATAIII VATPGRLLDHLQKGTVSLDAL NTLVMDEADRMLDMGFSDAID DVIRFAPASRQTLLFSATWPEAI AAISGRVQRDPLAIEIDSTDALP PIEQQFYETSSKGKIPLLQRTLV RFANGSARVLVATDVAARGLD IKSLELVVNFELAWDPEVHVHR IGRTARAGNSGLAISFCAPEEAQ RANIISDMLQIKLNWQTPPANS SIATLEAEMATLCIDGGKKAKM RPGDVLGALTGDIGLDGADIGK IAVHPAHVYVAVRQAVAHKA WKQLQGGKIKGKTCRRCYPLA APYVRHCSLVGRAASHQALWH VAGFGLYALFRTGGCRLRCR YSGGLRLIRPTVAAGICRPDKT RKRRIRHRAPFPDAARAPYPAY RSQYNQFFFTNSDFSFIGPKPSM LQSMLCAPCNQADVFHFGANV SASKSYLFTFRSLMMVTLSPCS QQIAVSIFNDQLVAFLCRFCVV
						PLRRHIHGQMYMPLSS
25112	55480	Α	25250	196	330	
25113	55481	A	25251	62	617	NRRLPSGGDEMSAQVSLELHH RI\SQF\LVQESSCWTDW\KFRD GVAQLRPKEVRF\TMRTTVNGQ TRVPRKGVQPPTTWIFNDTKDQ LERRIARLETGMAWAEEPPSRT RHLISNCQISETDIPNVFAVRVN YLLYRAQK\ERDETFYVGTRFD KVPPSE\DTTWRCLKRDIVLDQ AVITSHNLSVLF
25114	55482	Α	25252	1	318	
25115	55483	A	25253	1	510	
25116	55484	A	25254	9	79	
25117 25118	55485	A	25255 25256	863	1245	FSQIYLARSALSSAHSSRSPWQH SVWTLVRSVTNLETSANSALG NLWLRRATAHALKALSPNFWR LAHLSLCLADEAFPSSRS/IIELW QQAGSFISSLATFCKSSFIPFSKE KEFRGIKVRMSSGNFI
25119	55487	A	25257	909	1069	GILSPFLVFVRFVKDQIVVDVW YYF*GLCSVPLVDISVLVPVPCC FGYCSLVV
25120	55488	В	25258	1	1878	
25121	55489	Α	25259	778	990	
25122	55490	A	25260	312	635	
25123	55491	Α	25261	4047	4276	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25124	55492	A	25262	1599	11776	WFPASSMSQPKKAQDQTDSQP
25121	331,72	`	20202	1.577		NSTRGTERSWYHSF*NGSKQ*K
						KSNSSPTHFMRPASS
25125	55493	A	25263	1	162	
25126	55494	Α	25264	1	3370	MNRAPIPKQSKDKTSRETKNTS
						RSKTRERTDELEGLSTDNHAET
						QHAETNTDMTHERYTQIRNHT
						YKTTHQMEDKRTKPKREKEIH
						NRPDDMAKAGIPRSILNTGHPG
						KLMQTEVHHVRTLKIMLKVYS
		l				RALQEELQFSSKAIGRLFPCAD
! !						DLLETHSHFLAQTHERSGYVPL
						CVCATHTKFYEFMEDVNTAYT
		ĺ				KRHTMFRSLSMVSECLFSLING
						DDMFVTFAAMQAQQGRSSLV
						WLFSQLYLYSFISLFIYMV
25127	55495	В	25265	44	238	
25128	55496	Α	25266	1	444	MPAEFITG\ERS\SEGFF\RIHAGI
						EQAISRGLAYAPYADLKNLDD
						KTIASFQQQLSDMGYKFQFITL
						AGIHSMWFNMFDLANAYAQG
						EGMKHYVEKVQQPEFAAKKD
						GYTFVSHQQEVGTGYFDKVTTI
						IQGGTSSVTALTGSTEESQF
25129	55497	A	25267	710	1395	DLPGISSKWVIAVIFGHLIVGIFI
		1	ŀ			PPAIIVRAGIT*RQVR/WIAAHE
						GKAPQQIIIMAHLDTYAPLSDA
		1	į			DADANLGGLTLQGMDDNAAG
					;	LGVTLQLAERLKNTPTEYGIRF
					VATSGEEEGKLGAENLLKRMS	
						DTEKKNTLLVINLDNLIVGDKL
						YFNSGVQTLEARHQRSFNTVKP
						WPDSVISNRPFAGNTGSVDDND
						EIQRNLNDTNYNRMWEYNNRG
						VGSKVVAEAKK
25130	55498	Α	25268	2	2105	
25131	55499	Α	25269	754	1133	RAANQHQTVDLTRLQTRIAQR
	-					LFNRHSQAIEQRQPQIFQFVLVK
1		1				FAFPRFFLILPANVQWRLIATFC
						RFHTGVQGCQFTLVKRGPGDIR
					ŀ	DYQQTNLDWYNSRNTTKLDES
						VNRV/HAAIQS*RQNGL
25132	55500	В	25270	1	1275	
25133	55501	С	25271	1	1167	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25134	55502	Α	25272	275	2424	SPAMHFQSFT/WQNNLDDNTIA
						SFQQHLSDMGYKFQFITLAGIH
		Ì				SMWFNMFDLANAYAQGEGMK
						HYVEKVQQPEFAAAKDGYTFV
		İ				SHQQEVGTGVYAASGNSLLLM
		ŀ				RALNAAWPGIIDCSNHFARLRC
		ľ				SVWSIPRSDLRCAAAFRTGRLA
		ŀ				CCPAGDEKPYPSLRSSRWSGRG
		İ				ATALHYYTAKRCRPDKTRQRRI
		ļ				RQPARALDVGCGVTALSDLPG
						CRTSQRCRPDKTRQRRIRQPAR
						ALDVGCGVTALSDLRGAALHG
						SLANYRVVNFLFRLPDAMILSV
						LLRSCPFVFPLLRRLRAGAPVGP
						YCTGTPITPLPSTRLFAPLPNRC
						SMAGMYGGLQPVPHPAIGARC
	:					LMRCLTHLIRPTMGTGSMAEE
						ALGTSFKRLCAHYSRMRCWRIL
						SGLRGARPNKIVCRIRRLRRIRPI
ĺ						LRLRFPRHSRVVFIFGGGNPRR
			į			AKQRQTGEEVEKCHQPQHGTK
			ļ			CCARGGFAACLVFEYVVVFWT
						SLPDSFCRKSFPTRIRGSMVPSL
						WDFPPGVDGRKGKTSWNGVPG
						PCVTCGLGLTCPESPCCPGLSLG
						DGYPEKVSVGPPIPLKFQAILQA
						FHEIFGEGSCHIASVAVESGVPE
						QPFGSEETRAGARNRVANARR
						LLPEADFWVAIEAGIDGDSTFS
					:	WVVIENASQRGEARSATLPLPA
						VILEKVREGEALGPVMSRYTGI
						DEIGRKEGAIGVFTAGKLTRAS
25135	55503	Α	25273	1	441	
25136	55504	Α	25274	3	2919	AVSNCSEGLSKEWGPRPCRHR
ĺ						GMEKLPLGGKFWPTQ/EAVRT
						VRARLQAD/VTGVPTVLVARTD
						ADASDLITSDCDPYDSEFMTGE
						RTSEGFFRTHAGIEQAISRGLAY
						APYADLVWCETSTPDLELARRF
						AQAIHAKYPGKLLAYNCSPSFN
						WQKNLDDKTIASFQQQLSDMG
						YKFQFITLAGIHSMWFNMFDLA
						NAYAQGEGMKHYVEKVQQPE
		•				FAAAKDGYTFVSHQQEVGTGY
_						FDKVTTIIQGGTPDKAFT
25137	55505	A	25275	432	549	

SEQ ID NO:	SEQ ID NO: of peptide sequence	•	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25138	55506	A	25276	652	2689	KFSVGYAVQNFNATVSTKQFQ
						VFSEQNTSSYFAEPQLDVNYYQ
						NDVVHFVTTRDDMPEATRVHL
						EPTINLPLSNNWGSINTEAKLLA
						THYQQTNLDWYNSRNTTKLDE
						SVNRVMPQFKVDGRMVFERD
						MEMLAPGYTQTLEPRAQYLYV
						PYRDQSDIYNYDSSLLQSDYSG
						LFRDRTYGGLDRIASANQVTTG
		Ī				VTSRIYDDAAVERFNISVGQIY
						YFTESRTGDDNITWENDDKTGS
						LVWAGDTYWRISERWGLRGGI
						QYDTRLDNVATSNSSIEYRRDE
						DRLVQLNYRYASPEYIQATLPN
						VNHHITMEHLHMKTRTQQIEEL
						QKEWTQPRWEGIT\RPYSAEDV
						VKLRGSVNPECTLAQLGAAKM
						WRLLHGESKKGYINSLGALTG
						GQALQQAKAGIEAVYLSGWQV
						AADANLAASMYPDQSLYPANS
						VPAVVERINNTFRRADQIQWSA
						GIEPGDPRYVDYFLPIVADAEA
						GFGGVLNAFELMKAMIEAGAA
						AVHFEDQLASVKKMRHMGRT
						ASDQPWLAYAPYADLVWCEPP
						PPDLELARRFAQAIHAKYPGKL
						LAYNCSPSFNWQKNLDDKTIAS
						FQQQLSDMGYKFQFIT\LAGIHS
						MWFNMFDLANAYAQGEGMK
						HYVEKVQQPEFAAAKDGYTFV
		ĺ				SHQQEVGTGYFDKVTTIIQGGT
						SSVTALTGSTEESQF

SEQ ID		Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	noa	in USSN 09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25139	55507	A	25277	134	1695	FPDPARLPPVSVNHHITMEHLH MKTRTQQIEELQKEWTQPRWE
						GITRPYSAEDGGKLRGSVNPEC
						TLAQLGAAKMWRLLHGESKK
						GYINSLGALTGGQALQQAKAGI
	1					EAVYLSGWQVAADANLAASM
						YPDQSLYPANSVPAVVERINNT
ŀ						FRRADQI/QWSAGIEPGDPRYV
						DYFLPIVADAEAGFGGVLNAFE
						LMKAMIEAGAAAVHFEDQLAS
						VKKCGHMGGKVLVPTQEAIQK
						LVAARLAADVTGVPTLLVART
						DADAADLITSDCDPYDSEFITGE
						RTSEGFFRTHAGIEQAISRGLAY
	İ					APYADLVWCETSTPDLELARRF AQAIHAKYPGKLLAYNCSPSFN
						WQKNLDDKTIASFQQQLSDMG
						YKFQFITLAGIHSMWFNMFDLA
						KAYAQGEGMKHYVEKVQQPE
						FAAAKDGYTFVSHQQENRSSK
	•		}			QQQPLLTVGRIRRSRRIRQSVPR
						TGIGCASRRFNTAKPWADAVIS
						NRPFAGNTGSVDDNDEIQRNLN
						DTNYNRMWEYNNRGVGSKVV
						AEAKK
25140	55508	Α	25278	230	380	
25141	55509	Α	25279	432	1247	PSQEPSVHGRWMA*KGHSRIV*
						WRAYD*TH\FNPVLKKSVSLGS
						NIRHMLAEEATTEELDGT\GPV
						LV*PNRH*PAYWMRAYLNLKK
						MSCRSFRRKA*RW*CETQTVV
		i				HRDRVDQGNNHFDVVARHYH
						LYAFLRKERQLIFFKFKYARIQY
05140	155510	 	25200	242	120	DDDCVCVI ODI DCAVII DVV*D
25142	55510	A	25280	243	430	RRPSYSVLQRLPSAVLLPYY*R DWYHRTAGRRRDGNAGRQHQ
						NGCYPDPPDRDGRRSAFRNP
25143	55511	A	25281	10	638	SSVPTAGSRYNHLRAPRYSPLF
23143	33311	^	23201	10	038	ANYSMSDRCCKPFG/LDSSTG*
						WFQ*RSVPLPILLPDTSRHQL*C
			1			SDPTRNGPA\QACQRFFHIVERV
		1				ARVHHARFAATVTVDHVVIDQ
		1	ľ			RFFQICRT/PWCRLHRHHWRSSS
		1				GTR*TF*SPARHFCLLHNDSSYR
		1				STARPTCSGTDVPDRRCSDAHL
		1				CDHWSGSRAHADTDAECWCL
						NPQPAPSLLQCRR
25144	55512	Α	25282	803	1622	
25145	55513	A	25283	498	686	RRPSYSVLQRLPSAVLLPYY*R
			1			DWYHRTAGRRRDGNAGRQHQ
						NGCYPDPPDRDGRRSAFRNP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25146	55514	Α	25284	1	2063	MKNFMENVLRYLSDDKWKPD
						AKASMTVGTNLDTVYFKRHGQ
						VTGNSAAFDFHPDFAGISVEHL
		1				SSYGDLDPQEMPLLILNGFEYV
			,			TQVGNDPYAIPLRADTSKPKLT
						QQDVTDLIAYLNKGGSVLIMEN
						VMSNLKEESASGFVRLLDAAG
				į		LSMALNKSVVNNDPQGYPNRV
						RQQRATGIWVYERYPAVDGAL
						PYTIDSKTGEVKWKYHVENKP
						DDKPKLEVASWLEDVDGNRKR
						NFSNDARANGTAAFTDCETQT
]						VVHRDRVDQGHNHFDLVARH
						YHLYAFRQFDGTSHVSSTEVEL
1						RTVAFEERSMTAAFISWTECTL
						QIRTCPQNPTGKVWTCDELEIM
·		1				ADLCERHGVRVISDEIHMDMV
						WGEQPHIPWSNVARGDWALLT
						SGSKSFNIPALTGAYGIIENSSSR
		İ				DAYLSALKGRDGLSSPSVLALT
						AHIAAYQQGAPWLDALRIYLK
						DNLTYIADKMNAAFPELNWQIP
		1				QSTYLAWLDLRPLNIDDNALQ
		ŀ				KAL\IDQEKVAIMPGYTYGEEG
						RGFVRLNAGCPRSKLEKGVAG
						LINAIRAKKSATMIDTTLPLTDI
						HRHLDGNIRPQTILELGRQYNIS
						LPAQSLETLIPHVQVIANEPDLV
						SFLTKLDWGVKVLASLDACRR
						VAFENIEDAARHGLHYVELRFS
	ļ					PGYMAMAHQLPVAGVVEAVID
						GVREGCRTFGVQAKLIGIM
25147	55515	Α	25285	3	371	
25148	55516	Α	25286	63	180	
25149	55517	Α	25287	2	205	
25150	55518	Α	25288	495	707	REFSSHQKGLRDRPAHKAVLG
						DNQLFSGGTGAPVWPG/HSENV
						VPVDPT\MKRLPDDVNHPALDY
	<u> </u>					YWRLSFA

NO: of peptide sequence nod in USSN 09/540,217 location of first codon for peptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod feptide sequence nod nod feptide sequence nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod nod	n,
Sequence	
25151 55519 A 25289 1 1713 MTWAYSPSYGKPLSVPQ MTRFGMDVTLAHPEGYL VVEVAKNNAKASGGSFR MEEAFKDADIVYPKSWA MEERTELLRANDHEGVK HCL\AQNAQHK\DWHSIE EP\TREGEAQNKHGRP\AF SCKEGRSDQKAYDEKSR. PIDDNVVLNKGDVLQVS RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLIFAGIMLGFFN PTFGYIPQGALSMVKEFG MAGEIQVDCNTTYPHIWM GVLFAKQYLLRLGIILYG SQIADVGISGIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIG GVDVPNASLMIENPERL\ LHQLRGRVGRGAGLLT KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKU DHEGLKAVLERQCLAQN\ DWHCTEEMMELTRDGE\(TRUST)	rtion)
MTRFGMDVTLAHPEGYT VVEVAKNNAKASGGSFR MEEAFKDADIVYPKSWA MEERTELLRANDHEGVK HCL\AQNAQHK\DWH\SIE EP\TREGEAQNKHGRP\AR SCKEGRSDQKAYDEKSR PIDDNVVLNKGDVLQVS RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWR GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGGWR 25152 55520 A 25290 629 1054 ASGRIFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN DWHCTEEMMELTRDGE	
MTRFGMDVTLAHPEGYT VVEVAKNNAKASGGSFR MEEAFKDADIVYPKSWA MEERTELLRANDHEGVK HCL\AQNAQHK\DWH\SIE EP\TREGEAQNKHGRP\AR SCKEGRSDQKAYDEKSR PIDDNVVLNKGDVLQVS RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWR GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGGWR 25152 55520 A 25290 629 1054 ASGRIFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN DWHCTEEMMELTRDGE	HIGL
VVEVAKNNAKASGSFR MEEAFKDADIVYPKSWA MEERTELLRANDHEGVK HCL\AQNAQHK\DWH\SIE EP\TREGEAQNKHGRP\AB SCKEGRSDQKAYDEKSRI PIDDNVVLNKGDVLQVS RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL- LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/ DWHCTEEMMELTRDGE/	
MEEAFKDADIVYPKSWA MEERTELLRANDHEGVK HCL\AQNAQHK\DWH\SIE EP\TREGEAQNKHGRP\AE SCKEGRBQKAYDEKSR! PIDDNVVLNKGDVLQVSG RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFN PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWE GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGG GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKE YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN DWHCTEEMMELTRDGE	
MEERTELLRANDHEGVK HCL\AQNAQHK\DWH\SIE EP\TREGEAQNKHGRP\AE SCKEGRSDQKAYDEKSR PIDDNVVLNKGDVLQVSG RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGG GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKE YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/ DWHCTEEMMELTRDGE/	•
HCL\aQnaQhk\dwh\sie Ep\tregeaQnkhgrp\ab SCKEGRSQQKAYDEKSRI PIDDNVVLNKGDVLQVSG RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGG GVDVPNASLMIENPERL/LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAL GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI GAPLPLTANSDCNGYWR AAPLPLTANSDCNGYWR ASGRFREVTG/MEEAFKE YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/DWHCTEEMMELTRDGE/	
EP\TREGEAQNKHGRP\AI SCKEGRSDQKAYDEKSRI PIDDNVVLNKGDVLQVSG RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWM GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERLI LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKLD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNI DWHCTEEMMELTRDGEI	
SCKEGRSDQKAYDEKSRI PIDDNVVLNKGDVLQVSG RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWN GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERLI LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKLD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNI DWHCTEEMMELTRDGEI	
PIDDNVLNKGDVLQVSG RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKE YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/ DWHCTEEMMELTRDGE/	
RVKTIADRIGFISIHSQVTI CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKE YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/ DWHCTEEMMELTRDGE/	
CAFFVIGLMIGMITFQFST MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
MGNAAGLLFAGIMLGFM PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
PTFGYIPQGALSMVKEFG MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
MAGEIQVDGNTIYPHIWK GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
GVLFAKQYLLRLGIILYG SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERLE LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKE YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNE DWHCTEEMMELTRDGEE	
SQIADVGISGIIIDVLTLSS ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/ DWHCTEEMMELTRDGE/	
ACFLGQKVFGLDKHTSW GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERL LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
GSSICGAAAVLATEPVVK KVTVAVATVVIFGTVAIF YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERLE LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERLA LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
YPLMSQWFSPETFGIYIGS GVDVPNASLMIIENPERLA LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
GVDVPNASLMIIENPERLA LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	
LHQLRGRVGRGAGLLTA KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/ DWHCTEEMMELTRDGE/	
KRRFLKAAPEVTRTGLAI GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/ DWHCTEEMMELTRDGE/	
GLPLQTGYELASMQGLV LSPQGTDTFAMLDAFRAI AAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/ DWHCTEEMMELTRDGE/	
LSPQGTDTFAMLDAFRAMAAPLPLTANSDCNGYWR 25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKE YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQN/DWHCTEEMMELTRDGE/	
25152 55520 A 25290 629 1054 ASGRFREVTG/MEEAFKD YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNADWHCTEEMMELTRDGEA	
YPQSWAPYKVMEERTVL DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	RLAG
DHEGLKA\LEKQCLAQNA DWHCTEEMMELTRDGEA	ADI/V
DWHCTEEMMELTRDGE	LRAN
	QHK
HOWDADISGVSCKEGEV	LYM
I I I I I I I I I I I I I I I I I I I	EGVF
EKYRIATYKEASWKPYIL	\AMI
CPVNTPNQVHCSSNC	
25153 55521 A 25291 235 780	
25154 55522 A 25292 1 2783 MTYRNNHQTGVAFLFQT	
DGGVKTFDGYRVQTQSR	-
EVTDVEINLFCHPLCIIMQ	IFTV
HISEELATFIVRGFRFRRG	KAAI
GIFLVNNQLQPDIVHRGF	TEH
HDVRGIKHFTLVEHVITG	3GFG
NTRFTFFRAGNNKMPRLO	iVGA
GWAVLQGVPVRFGVWPI	PEYP
AYACGKFNGAIRAGDAA	QLAA
YAQAFIQLHGPIDAGDGV	NRA
DHRAGDFFEIVFQNLFAI	DVD
LAINHAVNAQLRLRNP	

SEV ID	ISEO ID NO-	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
SEQ ID NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25155	55523	A	25293	365	722	TSCLSALAIK/QQHLDEKKELG
		1				HLSAGARRLILGIIVTF/CADSG
						VNLRYSAV*PAGAVYLPDAAV
		1				GGSADRTADAGALLGANVDCA
		Ī				VADRFLPLYLVALHLYAELGRS
						GQPGVRAYSALR
25156	55524	Α	25294	798	1094	
25157	55525	Α	25295	1	2780	MGIVASASVQAAEIYNKDGNK
						LDVYGKVKAMHYMSDNASKD
						GDQSYIRFGFKGETQINDQLTG
						YGRWEAEFAAWTDMFPEFGG
						DSSAQTDNFMTKRASGLATYR
						KTTSSAYRWPELNLQYHRCTS
		1				AARAMLLSGRPVMQINERTAV
						RRQMTVYLRVERIHEIDPLVSL
						SHSMSVIPIDVERPIAAAPCTPG
						LWRTKTPLRVPSTYPLRLPGRA
						RVVCHRTFRLHLCKDWVFMFS
25150	55506	ļ	25206		200	GLLIILVPLIVGYLIPLRQQA
25158	55526	Α	25296	66	300	QSTYCDNPPVPVSRGL*P*RRSS
		i				STSSAGYQK*RYCQHRHDPASR
						IRTVQRTDSQRTVKRKFMLLVP EASVPASEIC
25150	55527	 	25297	240	424	RCSRDEHDFQLIANLRIVVITFA
25159	33321	Α	23297	240	424	TFA*CTA*LPLGNHSVVAAH*A
				1		G*GGAPVTAQVFVQHG
25160	55528	A	25298	1	1425	O GOAL VIAQVI VQIIG
25161	55529	A	25299	1	471	
25162	55530	A	25300	1269	1390	
25163	55531	Α	25301	601	1119	VAAGAV*RGH*NVQLGKLFFV
						ASDLRINAIAGGVVVFS*FIQGL
		1				TNAIGSGFNNARCRKPTGQVA
		1				ADGGRDAGFFTIANNQCTYAA
						GIFFQCSGEFVGIQFQFAIAGIGR
						DTNLRQRFIAGRTDGITQHRFQ
						LRLNRPQRGVAEQCTIARTLAIS
ļ						SRMAVYLVRLLKVSISEL
25164	55532	A	25302	1666	1769	RLKRHPPLRPSWPPY*TGAHVW
1						HFSPLANWKPGI
25165	55533	Α	25303	1	1970	MKDDIFFRTSGGGVTLSGGEVL
						MQAEFATRFLQRLRLWGVSCA
						IETAGDAPASKLLPLAKLCDEV
						LFDLKIMDATQARDVVKMNLP
						RVLENLRLLVSEGVNVIPRLPLI
						PGFTLSRENMQQALDVLIPLNIR
						QIHLLPFHQYGEPKYRLLGKTW
						SMKEVPAPSSADVATMREMAE
						RAGLQVTVGEKWGVSIETQGA
						LGTENRLADEDIRRADVALLIT
						DIELAGAERFEHCRYVQCSIYA
						FLREPQRVMSAVREV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25166	55534	Α	25304	2804	3494	LSLAEWKCLNPSQRALYREVM
		ŀ				LENYRNLEAVGQLQPQSTQRIP
						*GRICNGDRLTSQKGLDL\VLEA
			İ			LPGLL\EQGGQLALLGAGDPVL
		ł				QEGFLAAAAEYPGQVGVQIGY
						HEAFSHRIMGGADVILVPNRFE
						PCGLTQLYGLKYGTLPLVRRTG
						GLADTVSDCSLENLADGVASG
		ſ				FVFEDSNAWSLLRAIRRAFVLW
						SRPSLWRFVQRQAMAMDFSW
						Q\VAAKSYRELYYRLK
25167	55535	Α	25305	1	1306	QTT. III. III.
25168	55536	A	25306	818	1658	
25169	55537	Α	25307	2611	2826	
25170	55538	Α	25308	609	2359	PPPKIAPVWGCRKKSLPARSKSI
			,			FCFALFFPFFLNGEIILPESPCPF
						WTLFRNDGGKKRDPLAPVYLL
						GKLTISPGPICPDMIGVLFDSTV
	2					RREVAHIGNVQHRFCGPLFLKL
						VEFIDFILTIDIAAIIRQYLVVICE
						VDQRINQIAIATRIFRTGYTAAD
						LRQHLMQLLIFLVVFTRFVALT
						TQFFHFFRCVTKNKDIVSTDMI
					:	QHLDIRTVQRTDSQRTVKRKLH
						VTGAGSFRPCQRNLFRQICRRN
						DQLRQANAVVRDEHDFQLIAN
						LRIVVDHFRYVIDQMNNVLRH
						VIGRSRLTAKNIHARYPLRIRVG
						LDAVIAEMASGLDPFWRPDVV
						HAHDWHAGLAPAYLAARGRP
						AKSEFTVHNLAYQGMFYAHH
						MNDIQLPWSFFNIHGLEFNGQIS
						FLKAGLYYADHITAVSPTYARE
						ITEPQFAYGMEGLLQQRHRERR
						LFGVLNGVDEKIWSPETDLLLA
						SRYTRDTLEDKAENKRQLQNA
						MGLKVDDKVPLFAVVSRLTSQ
			-	}		KGLDLVLEALPGLLEQGGQLA
						LLGAGDPVLQEGFLAAAAEYP
						GQVGVQIGYHEAFSHRIMGGA
						DVILVPSRFEPCGLTQLYGLKY
						GTLPLVRRTGGLADTVSDCSLE
						NLADGVASGFVFEDSNAWSLL
						RAIRRAFVL/WSRAGSWRYSAR
						AIRCCRKVSLRRQRNTPVRWAF
						RLAITKHFRIALWAARTSFWCP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	•	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			:	sequence		
25171	55539	A	25309	1028	1632	TSASAGTARNFSSNFAVRTTGH
23171			23303	1020	1332	STSEATSSSRLSLKSALPPILRAA
						SSASALMFFVIGDDFPTSRRISG
		j]		Y*SALSMVNSGSPIKRWPRITRS
		ĺ				D*MPRMVAGMTLSP/MQQGHG
						VHRTHEVHVVAPQRISFGIGSF
						AREVATIFGSSASVFRR*YRNH
						APNLLTLCSVCLQRRTPGQSPG
						RVFRFHDPVAFRQVSYFQRQTP
ļ						IA
25172	55540	Α	25310	1	336	
25173	55541	A	25311	1889	2257	SAMTFPRSRRISGY*SALSMVN
20175						SGSPIKRWPRITRSD*MPRMVA
						GMTLSPSSRVTVCTGRTKCTSL
						APQRISFGIGSFAREVATRRPYR
E						DIKEGFSQREAALLPQSLRVPQ
						TGTWSAAGAQA
25174	55542	В	25312	1	735	
25175	55543	Α	25313	1439	1744	SAMTFPRSRRISGY*SALSMVN
						SGSPIKRWPRITRSD*MPRMVA
						GMTLSPMQQGHVPWQLMVVL
						WNTRQRWKPSSPDRYVNRAMS
	٠.					LPSRKGEMLKPAPCRK
25176	55544	Α	25314	1209	1972	SAMTFPRSRRISGY*SALSMVN
						SGSPIKRWPRITRSD*MPRMVA
		1				GMTLSPSSRVTVCTGRTKCTSL
1						APQRISFGIGSFAREVATIFGSSA
1	:					SV/HFALHMGAIQQPFAFIGSQT
						LSLVNGNTATT\PQPSAALLGLP
1	1					SASNAWAIAGPRFSISRSGCASA
			1			RLATFSARRRGASKPLNGCLLR
1			1			ATLSSPDKFQQGSTGGMLWGV
				-		LRHSPKFSITADGREFTLHWPN
						TGAMTGCATPGEYVWYSTFDN
						TQRKYPHPWEL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	Ì	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25177	55545	Α	25315	1	1521	METQHRLMRTFQPRQQDWTY
25177	33343	l'`	25515	1	1321	ALWGGVKLRWLFFFLWGLFLF
						AGEEELGIRQTRTNDFLVTGDN
						LLRIFRFDVGNEDKVRQQFAVV
						RIHREVFLVTFHGVNQRFSRHR
						EEFLFEFCVTPKLNEGTTQRTN
						NRGIAVLNTGHRHPDLVAAVE
						QQLQQFTHTAYQIVPYESYVTL
						AEKINALAPVSGQAKTA\FFHP
						GSETG\ENGVKMVRAHTGTPG
						V\IAFRGG\FHGRTYMTMALTG
						KVAPYKIGFGPFP\GSVYHVPYP
						SDLHGISTQDSL\DAIERLFKSDI
		l				EAKQVAAIIFEPVQGEGGFNVA
		1				PKELVAAIRRLCDEHGIVMIAD
						EVQSGFARTGKLFAMDHYADK
						PDLMTMAKSLAGGMPLSGVVG
						NANIMDAPAPGGLGGTYAGNP
						LAVAAAHAVLNIIDKESLCERA
						NQLGQRLKNTLIDAKESVPAIA
						AVRGLGSMIAVEFNDPQTGEPS
						AAIAQKIQQRALAQGLLLLTCG
						AYGNVIRFLYPLTIPDAQFDAA
25178	55546	Α	25316	114	187	RSTISSVRYGDDAFKP*RQAWN
						VL
25179	55547	Α	25317	105	317	YLFFLSRFNHWISMEKKLGLSA
						LTALVLSS\SWARVFSVCRKIW
						RQLPARQHCSSAGVLLALAFYC
						WPLPC
25180	55548	A	25318	1	593	
25181	55549	A	25319	1	729	
25182	55550	A	25320	1	1005	
25183	55551	A	25321	1	1361	
25184	55552	В	25322	1	2793	
25185	55553	Α	25323	27	255	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25186	55554	Α	25324	225	1408	RIIYMHMESQKIWFALSTPMEIR
						NECCLPSHSSPKMYLGACLF\PL
						SSSWGIDDRDDLLRTIHRMIDN
					Ì	GHAARLAGFYHRWFRYSPCEW
						RDYLAELNEQGQAYAQFVAST
						AECCGEGGIKAWDYVRMGFLS
						RMGVLNNWLSEEESLWIQSRIH
						LRALRYYSNWRQYFAGYTFGR
ŀ				ĺ		QYWQSPEDDHLPLLREFLAQET
						TDVDIIRRAYLALLPSFHTETDP
						QGFKQLRHAYDDALPIAQSPAK
						SVLQPEEYEWEALLVEMLGLL
						HRIAMVQLSPAALGNDMAAIE
						LRMRELARTIPPTDIQLYYQTLL
		ĺ				IGRKELPYRTCQLHACTSADTP
		ļ				LRKIPRFSGKLHFYAFCPNIYSS
						YHSAAKYGGSCQPGSTAHRLG
						YYWRWHFIAGLCHADPHAHSS
25187	55555	Α	25325	230	380	
25188	55556	Α	25326	432	1247	PSQEPSVHGRWMA*KGHSRIV*
						WRAYD*TH\FNPVLKKSVSLGS
						NIRHMLAEEATTEELDGT\GPV
					İ	LV*PNRH*PAYWMRAYLNLKK
						MSCRSFRRKA*RW*CETQTVV
						HRDRVDQGNNHFDVVARHYH
						LYAFLRKERQLIFFKFKYARIQY
						AG
25189	55557	Α	25327	10	638	SSVPTAGSRYNHLRAPRYSPLF
						ANYSMSDRCCKPFG/LDSSTG*
						WFQ*RSVPLPILLPDTSRHQL*C
						SDPTRNGPA\QACQRFFHIVERV
						ARVHHARFAATVTVDHVVIDQ
		ŀ				RFFQICRT/PWCRLHRHHWRSSS
						GTR*TF*SPARHFCLLHNDSSYR
						STARPTCSGTDVPDRRCSDAHL
•						CDHWSGSRAHADTDAECWCL
25100	65550	_	05220	002	1622	NPQPAPSLLQCRR
25190	55558	⊢	25328	803	1622	PPDOVOVI ON POATE PARTE
25191	55559	Α	25329	498	686	RRPSYSVLQRLPSAVLLPYY*R
						DWYHRTAGRRRDGNAGRQHQ
	<u>L</u>	<u> </u>				NGCYPDPPDRDGRRSAFRNP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
25102	155560	1.	105220	1.	2063	 MKNFMENVLRYLSDDKWKPD
25192	55560	Α	25330	1	2003	t ·
						AKASMTVGTNLDTVYFKRHGQ
						VTGNSAAFDFHPDFAGISVEHL
				ļ		SSYGDLDPQEMPLLILNGFEYV
						TQVGNDPYAIPLRADTSKPKLT
-			}			QQDVTDLIAYLNKGGSVLIMEN
		ł				VMSNLKEESASGFVRLLDAAG
		ŀ				LSMALNKSVVNNDPQGYPNRV
		ĺ				RQQRATGIWVYERYPAVDGAL
1				İ		PYTIDSKTGEVKWKYHVENKP
-						DDKPKLEVASWLEDVDGNRKR
						NFSNDARANGTAAFTDCETQT
						VVHRDRVDQGHNHFDLVARH
						YHLYAFRQFDGTSHVSSTEVEL
						RTVAFEERSMTAAFISWTECTL
						QIRTCPQNPTGKVWTCDELEIM
						ADLCERHGVRVISDEIHMDMV
						WGEQPHIPWSNVARGDWALLT
						SGSKSFNIPALTGAYGIIENSSSR
						DAYLSALKGRDGLSSPSVLALT
						AHIAAYQQGAPWLDALRIYLK
						DNLTYIADKMNAAFPELNWQIP
						QSTYLAWLDLRPLNIDDNALQ
}		ļ				KAL\IDQEKVAIMPGYTYGEEG
						RGFVRLNAGCPRSKLEKGVAG
1		ĺ				LINAIRAKKSATMIDTTLPLTDI
						HRHLDGNIRPQTILELGRQYNIS
		İ				LPAQSLETLIPHVQVIANEPDLV
				:		SFLTKLDWGVKVLASLDACRR
						VAFENIEDAARHGLHYVELRFS
						PGYMAMAHQLPVAGVVEAVID
İ						GVREGCRTFGVOAKLIGIM
25193	55561	A	25331	2	696	FLLLLWEIRKSSQQTTPTEMNA
				<u> </u>		AEMAQLVAGVDEVGRGPLVG
						AVVTAAVILDPARPIAGLNDSK
		1				KLSEK\RRLALYEEIKEKALSWS
		į				LGRAEPPEIDELNILHATMLAM
ŀ			}			QRAVVGLHIAPEYVLIDGNRCP
						KLPMPAMAVVKGDSRVPEISA
						ASILAKVTRDAEMAALDIVFPQ
						YGFAQHKGYPTAFHLEKLAEH
						GATEHHRRSFGPVKRA/LGTCV
						LILVSRLSKPESEDVL
25194	55562	A	25332	2	63	LIL V SINLSINF ESEU V L
25194	55563	В	25333	138	182	
25196	55564	A	25334	356	706	ISDICAPNMVSAIHAIPYTATMA
	1	`			- 30	IIPERKTSFFEIDAGLVLVRIIISA
		1				PVSSINI*ITSTIDSGLPPKDGRRS
						ESAPRRATTAIAPRKIHIASQLL
						VVIIIRSLVGCRMRRWRVLSD
25197	55565	A	25335	1746	1873	T TIMBLY GORWING WICY LSD
23171	122202	1,,		11.10	1.07.5	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25198	55566	A	25336	11998	3626	<u> </u> PKAARMANVPNNTTGTAMVGI
23190	33300		23330	1770	5020	SVA/RHILQEQIHHQEHQHNGFE
ļ	1					QRFYHFVNRKLDEWRGVIRVE
						NLHS/AAGRMAVIPPVLP*WLS
		1				RYREHSRRLPA*SPKGEAASNP
						EVIARTLRKLKNQPELSEDTIKK
						AVESLSLELVLTAHPTEITRRTL
1		ŀ				IHKMVEVNACLKQLDNKDIAD
						YEHNQLMRRLRQLIAQSWHTD
						EIRKLRPSPVDEAKWGLCRNGK
						TACGKAYQITWAELNKQLEEK
	<u> </u>	!				PRLQTCPSKFGSGPFYF
25199	55567	Α	25337	243	604	PRFPFPLPPATSSVQTAMTDG*N
						ECERICVGYLRYSSHSAPDARR
						QRARTHVPVGKHALPALTKAT
						HPSWWWQCAAMAEDSTDARQ
						LKA*WGVNGFLDLRSATMDGR
						LYPVAMCLLPATK
25200	55568	Α	25338	1	3287	MPITERRSVEKEQLIEIANTIMPF
						GKYKGRRLIDLPEEYLLWFARK
						DEFPAEARYTHIDLHVPAPLLV
15		1				MFTLDTRQSACDRNPHLPIHVC
						SSWTSPKCAPRLWTRALLPRTR
		1				TPPYATGTTIGSRASWFVTFPD
		1				ASFYRRKAPRQEFHTSLQGRSL
		ł				YVRQRSTANPTEITRRTLIHKM
		1			,	VEVNACLKQLDNKDIADYEHN
	İ					QLMRRLRQLIAQSWHTDEIRKL
1						RPSPEAIMTEQEKTSAVVEETR
						EAVDTTSQPVAT
25201	55569	A	25339	1	1425	
25202	55570	Α	25340	534	1391	RQKRRLYQTYWYKQQPRYLPP
						EHRQRHRHQSAGVSRPYSGYG
						EVRPLSQYQPEWQCWEPDVEW
		1				VRQPPQDAITDPDFFCFYQPGM
						TFEQFVREFAEWFSQKRPAAM
						MIGIRADESYNRFVAIASLNKQ
						RFADDKPWTGGHSWYIYPIYD
		1				WKVADIWTWYANHQSLCNPL
				•		YNLMYQAGVPLRHMRICEPFG
		1				PEQRQGL\WLYHVIEPDRWAA
		Ì				MCAR\VS\GYTTGGLYPGQARG
						AAV*TEALKPSGLPQVCALNQC
			Ì			LLSVHRYAGHVRQERCDGADG
						RPAVHERRQLL
25203	55571	A	25341	786	1009	HHSHEQQFQWPVGASGVDRW
						AVYLHQICPLTQFQY/WCRARQ
						EGIMSSPGQQVGFIA*VTSSFSP
						R*KGRRVSSA
25204	55572	A	25342	2446	2970	
25205	55573	A	25343	1	2184	
25203	55571	A	25341	786	1009	RPAVHERRQLL HHSHEQQFQWPVGASC AVYLHQICPLTQFQY/W EGIMSSPGQQVGFIA*V
-	55572	A	25342	2446	2970	
25205	55573	A	25343	1	2184	

SEQ ID	SEO ID NO:	Met	SEQ ID NO	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25206	55574	I _A	25344	706	1293	IHRLLIRRIHQIIDLRIWORTVEI
23206	33374	A	23344	700	1293	` ` `
						HGAPVRFIQMAAWTHIAVLLT*
	1					QLRQLRAGFQRNSWAIFYFFHH
						RKQYIFHFDLRRFPVIMHALFC
1						AVQRCADLMAHRRQKRRFCLA
						GFFCFRPCLSVFYHQFLQFSCTR
		1				FDLRRQLFAVAIQCMNIAFAAA
		1				LVDFDLPDIDGITLARQLAQQY
						PSLVLIGFSAHVIDETLQ
25207	55575	Α	25345	1	3291	
25208	55576	Α	25346	882	1019	
25209	55577	Α	25347	1	2265	
25210	55578	Α	25348	284	535	CVAASVTGTRVLIPPLLNNRSR
						RSPVRYVSKGNWWGSVCNCAS
		1				NNSNSVSR*SLPPMRSHAWRKV
		<u>L_</u>				RRIMRQLPLERPRPGFTI
25211	55579	Α	25349	131	208	LVRWLLSPGW*WKISPIISICLFP
						V
25212	55580	Α	25350	1	3108	MDSQDKYFEATQTVYEWCGV
	1					ATQLLAAYILLFDEYNEKKASA
						QKDILIKVLDDGITKLNEAQKS
						LLVSSQSFNNASGKLLALDSQL
						TNDFSEKSSYFQSQVDKIRKEA
						YAGAAAGVVAGPFGLIISYSIA
						AGVVEGKLIPELKNKLKSVQNF
						FTTLSNTVKQANKDIDAAKLKL
						TTEIAAIGEIKTETETTRFYVDY
						DDLMLSLLKEAAKKMINTCNE
						YQKRHGKKTLFETYLTDHPVTF
						RERLOMSIYKIPLPL
25213	55581	A	25351	1	296	
25214	55582	Α	25352	3	164	VGD/NVGTTARWGIFALIIVETL
						AWFTVVASLFAQHKQRAENLI
						QRLHLLWSFNP
25215	55583	В	25353	91	990	
25216	55584	Α	25354	1146	1881	RCGVQ\WTMYIRAVFVNLTNP
						KSIVFLAALFPQFIMPQQPQLM
						QYIVLGVTTIVVDIIVMIGYATL
						AQRIALWIKGPKQMKALNKIFG
						SLFMLHNAFAWFYLGAFLLTVI
		1				AWFLLGNAQKMPQTTLQWGIL
						VFLGVVASGIGYFMWNYGATO
						VDAGTLGIMNNMHVSAGLLVN
1						LAIWHQQPHWPTFITGALVILA
						SLWVHPNKRFRNWLTLLAIAH
						GGVQSAPQAFAQGFGHHQTLA
						LKTLYYPMR
25217	55585	A	25355	1943	2287	LK I L I I FIVIK
25217	55586	A	25356	44	274	
23210	72220	17	22330	177	1	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25219	55587	A	25357	876	1400	YRHVRDSPGNDRLFETWRSAG
						CAARLLRQSGETRLLSRVVRDY
						GGRWPVSLLDGLPDATWCTEK
					-	RGGFCTCATGRAGEKWAQFPE\
						SLLTNLANPKAIIYFGSVFSLFV
						GDNVGTTARWGIFALIIVETLA
						WFTVVASLFALPQMRRGYQRL
						AKWIDGFAGALFAGFGIHLIISR
25220	55588	A	25358	1	1428	
25221	55589	A	25359	421	797	IIFPFLLFHLCNQLSHLLQYAVV
20221		-				NAPVRRILPIQLLWQQLIQPGEK
						LKSIRRAIQHIGQTQIIGMFIQMP
						IMIAYETDRATGDGI/WAAH*T
						HEYPCHVRQSQFHENHDDVQE
						NPPHPWKIQTPGH
25222	55590	A	25360	1	1320	
25223	55591	A	25361	1	1251	MLLSIITVAFRNLEGIVKTHASL
23223	33371	1.	20001	1		AHLAQVEDISFEWIVVDGGSND
						GTREYLENLNGIFNLRFVSEPD
						NGIYDAMNKGIAMAQGKFALF
						LNSGDIFHQNAANFVRKLKMQ
						KDNVMITGDALLDFGDGHKIK
						RSANRAGLFLIGYGAFRIIVEFF
						RQPDAQFTGAWVQYISMGQILS
		-				IPMIVAGTMKQYLELMQKVLD
						EGTQKNDRTGTGTLSIFGHQMR
						FNLQDGFPLVTTKRCHLRSIIHE
ŀ						LLWFLQGDTNIAYLHENNVTI
		İ				WDEWADENGDLGPVY\ITTVL
					•	NQLKNDPDSRRIIVSAWNVGE/
				İ		LDKMALAPCHAFFQFYVPDGK
		1				LSCQLYQRSCDVFLGLPFNIAS
1						YALLVHMMAQQCDLEVGDFV
		1				WTGGDTHLYSNHMDQTHLQLS
						REPRPLPKLIIKRKPESIFDYRFE
25224	55592	A	25362	3	1327	REFREINGE EST DTRE
23224	22392	A	23302		1321	

SEQ ID NO:	SEQ ID NO: of peptide sequence		_	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25225	55593	A	25363		2443	MGLIGFVLTALWIKLIHNPTDH PRMSAEELKFISENGAVVDMD HKKPGSAAASGPKLHYIKQLLS NRMMLGVFFGQYFINTITWFFL TWFPIYLVQEKGMSILKVGLVA SIPALCGFAGGVLGGVFSDYLI KRGLSLTLARKLPIVLGMLLAS TIILCNYTNNTTLVVMLMALAF FGKGFGALGWPVISDTAPKEIV GLCGGVFNVFGNVASIVTPLVI GYLVSELHSFNAALVFVGCSAL MTMSARSATGHECVFWCNGKI RRLAMRLIYWCAAAPTGDVVC KLTGKWALIANSHRAGQPTRN VNEEVGQRLLEEMEWYDNYL NMGKTDRSANPSPGNKKGGLA NVVEKALGSIAKSGKSAIVEVL SPGQRPTKRGLIYAATPASDFV CGTQQVASGITVQVFTTGRGTP YGLMAVP/SH*NGNWKITASVT SIQPNTNELPNCCMNGVSTANT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25226	55594	Α	25364	1	2701	TFFLPAGEILPPRLEPQITLPVGS
						RRIISIAVTDLPEPESPHLRSGLG
		1				WASLCCFMVICKFFPAAEGLIL
						AAVCDSRGSRYWCNGNRRYG
		ł				YASDLLVRCGATVMFSEVTEV
						RDAIHLLTPRAVNEEVGKRLLE
						EMEWYDNYLNMGKTDRSANP
						SPGNKKGGLGKRGSSKALWLP
						LLQIQARKQVGLRVYTVACVL
]						RTGTWYDRNGPDWRVPGQLN
						MGNPAPETGRYRLAHPATPHD
						PRTTPDHPSHIKPHAPTNQQPDT
						LTPTHHHHTDTRDPSPPPPRTRT
						PRQSTHTTPPHPTTTTRHTDIHP
						PTHTNPPPPTRVLTRQLHHTNRI
						IPHRTHTTTCRRNHHNQMNTTT
						PPPPHTPQITHRHHNTTLREQTQ
						КНКОНІНСТОННТТККРРТРТТ
		1				RQYNTSIVGTIGGG\MVERKVIE
						ESLQALQERKPRLFHGRMARN
						GADAVGSDCGGAMSVFISVHG
		1				MRPRLVLIGAGHVNRAIAQSAA
		1				LLGFDIAVADIYRESLNPELFPP
		Ì				STTLLHAESFGAAVEALDIRPD
		1		1		NFVLIATNNQDREALDKLIEQPI
						AWLGLLASRRKVQLFLRQLRE
						KGVAEEHIARLHAPVGYNIGAE
						TPQEIAIRVLAEILQVKNNAPGG
						LMMKPSHPSGHQMVVIRGAGD
						IASGVALRLYHAGFNVIMLEVE
			-			KPTVIRCTVAFAQAVFDGEMT
						VEGVTARLATSSAEAMKLTER

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25227	55595	Α	25365	1	1262	MGNWHLSPCPAPAPGHSCLFL
						VGTSPRILAVPGHTGSKAINKA
						ALLLESYMSQALKNLLTLLNLE
						KIEEGLFRGQSEDLGLRQVFAG
	,					QVGG\QPLYAAKEPVPEERLVH
						SFHSYFLRPGDSKKPIIYDVETL
						RDVAPVLDVAGATPPTIRHQRV
						TALVAPKNGSRDVRSCAGKMP
						TDGSIRAVLLPFQVSTIGGVTYE
						RVGTKDQGPQCPHVQRSSAVR
						PGPEAAPGDNSYPPYEDRSTCG
						MQAPDASYSANPDLSFTWSPCL
						SEDALHTSPLYCTTPTSGGEAF
						KGLASDLVPHEDKMVLQNGKC
						FTSPRHTEWAVLRAGHALARG
			ŀ			RNLLDLAAGGMAWVSMSQEC
						PGPGTICSRDLQGTECLRMIQIL PPKDPVTRGSTHCNAGTKRELE
						EFSSQVTLTALIQPLMQVSAQF
						RAADGAP
25228	55596	Α	25366	2	176	RAADUAF
25229	55597	A	25367	1	1395	MRESLELPRDLLNCCDQNANS
23229	33371		25507	1	1373	DMDMKSRLRGYQMEIMNLLG
İ						AGVKDFELESDDLVYLVEEISK
						QQSIHDVAWLLLVAFVYICEQR
						NDLKSELIFTREAEHKSLENLEP
						DHVPEELDTSARNAGALTRRRE
						IRDAATLLRLGLAYGPGGMSLR
	·			•		EVTAWAQLHDVATLSDVALLK
1						RLRNAADWFGILAAQTLAVRA
1		1				AVTGCTSGKRLRLVDGTAISAP
İ						GGGSAEWRLHMGYDPHTCQFT
1						DFELTDSRDAERLDRFAQTADE
Ì						IRIADRGFGSRPECIRSLAFGEA
1						DYIVRVHWRGLRWLTAEGMRF
						DMMGFLRGLDCGKNGETTVMI
1						GNSGNKKAGAPFPARLIAVSLP
						PEKALISKTRLLSENRRKGRVV
						QAETLEAAGHVLLLTSLPEDEY
						SAEQVADCYRLRWQIELAFKR
				ŀ		LKSLLHLDALRAKEPELAKAWI
						FANLLAAFLIDDII\SHRWISPPE
25230	55598	Α	25368	482	765	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25231	55599	Α	25369	1	1575	MGSVTKSRGPTDSSAFFQINRD
						YSFLDYILGGCQLMFTPWARLL
						QSTCCEQCSKKAHDQFLADLA
						SILPSNTTPLIVSDAGFKVPWYK
						SVEKLGWYWLSRVRGKVQYA
						DLGAENWKPISNLHDMSSSHSK
						TLGYKSLTISNPISGQILLYKSRS
						KGRKNQRSTRTHCHHPSPKIYS
						ASAKEPWILATNLPVEIRTPKQL
						VNIYSKRMQIEETFRDLKSPAY
						GLGLRHSRTSSSERFDIMLLIAL
						MLQLTCWLAGVHAQKQAFRF
						MRLVDGTAISAPGGGSAEWRL
						HMGYDPHTCQFTDFELTDSRD
						AERLDRFAQTADEIRIADRGFG
						SRPECIRSLAFGEADYIVRVHW
						RGLRWLTAEGMRFDMMGFLR
						GLDCGKNGETTVMIGNSGNKK
						AGAPFPARLIAVSLPPEKALISK
						TRLLSENRRKGRVVQAETLEA
						AGHVLLLTSLPEDEYSAEQVAD
						CYRLRWQIELAFKRLKSLLHLD
						ALRAKEPELAKAWIFANLLAAF
0.7000	55500	ļ.,	25250		1044	LIDDII\SHRWISPPEVPDPKRRT
25232	55600	Α	25370		1341	MAGGRKFPDGGDRTTLCGLSQ
						PGVLAPRSPEKRRAINQRYPSK
						TGQLPASITDMMRTARHHLSG
						NGISVPKGAELAHPSTVLELFAI
						SAKPIHPEELDTSARNAGALTR
	ļ					RREIRDAATLLRLGLAYGPGG
						MSLREVTAWAQLHDVATLSDV
						ALLKRLRNAADWFGILAAQTL AVRAAVTGCTSGKRLRLVDGT
						AISAPGGGSAEWRLHMGYDPH
		ŀ				TCQFTDFELTDSRDAERLDRFA
						QTADEIRIADRGFGSRPECIRSL
						AFGEADYIVRVHWRGLRWLTA
		ĺ				EGMRFDMMGFLRGLDCGKNG
				•		ETTVMIGNSGNKKAGAPFPARL
1						IAVSLPPEKALISKTRLLSENRR
1						KGRVVQAETLEAAGHVLLLTS
1						LPEDEYSAEQVADCYRLRWQIE
						LAFKRLKSLLHLDALRAKEPEL
]						AKAWIFANLLAAFLIDDII\SHR
						WISPPEVPDPKRRT
25233	55601	В	25371	1	1233	
	122201		200,1	·	1.200	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25234	55602	Α	25372	1	2197	SHSRGSPGKAVDFTLVSHAAW
2323	33002	, `	23372	•	2197	FGLPHFSTPAFNGQAMMLIAPV
						AVILVAENLGHLKAVAGMTGR
						NMDPYMGRAFVGDGLATMLS
						GSVGGSGVTTYAENIGVMAVT
l						KVYSTLVFVAAAVIAMLLGFSP
						KFGALIHTIPAAVIGGASIVVFG
						LIAVAGARIWVQNRVDLSQNG
			,			CNPAYAEHHQHPGYSDKLKTR
						VSFSCLRIISTNLRRECRAQFRF
						HVDASARHGFRLWQIQGVAQP
						VFIITIREVEAAMRATAFGTEVG
						1
						RNSRCIRRFHQVIQFQALNALG
		1				VELAGIDSRDPRLKVLEEMKRT
						FHWGRQTISKHIKIWLGLMLGR AMTPLMLDATSGKLTVWDGA
		1				i i
						HAGAAMGILAVTADQNSAELA
						YYKSGSFRIEDVLWPSAVTDDN
						IKRNAFAADGSVNSYSVPFSSV
						PLLQRQGRIKYAVTLAKYRTNS
						NEQQESKFAQATLQWGGPWGT
						TWYGGGQYAEYYRAAMFGLG
						FNLGDFGAISFDATQAKSTLAD
						QSEHKGQSYRFLYAKTLNHLG
						TNFQLMGYRYSTSGFYTLSDT
						MYKHMDGYEFNDGDDEDTPM
						WSRYYNLLYTKRGKLQSLISKT
1						RLLSENRRKGRVVQAETLEAA
						GHVLLLTSLPEDEYSAEQVADC
						YRLRWQIELAFKRLKSLLHLDA
						LRAKEPELAKAWIFANLLAAFL
						IDDII\SHRWISPPERIPSARKWT
25235	55603	Α	25373	1	2754	
25236	55604	Α	25374	1	652	
25237	55605	Α	25375	1	1800	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	!	
25238	55606	Α	25376	1	2187	MAGPRYPVSVQGAALVQIKRL
						QTFAFSVRWSDGSDTFVRRSW
						DEFRQLKKTLKETFPVEAGLLR
		Ì				RSDRVLPKLLGQASLDAPLLGR
						VGRTSRGLARLQLLETYSRRLL
						ATAERVARSPTITGFFAPQPLDL
		ŀ				EPALPPGSRVILPTPEEQPLSRA
						AGRLSIHSLEAQSLRCLQPFCTQ
		Ì				DTRDRPFQAQAQESLDVLLRHP
						SGWWLVENEDRQTAWFPAPYL
						EEAAPGQGREGGPSLGSSGPQF
						CASRAYESSRADELSVPAGARV
						RVLETSDRGWWLCRYAGAGPE
						ELDTSARNAGALTRRREIRDAA
						TLLRLGLAYGPGGMSLREVTA
						WAQLHDVATLSDVALLKRLRN
						AADWFGILAAQTLAVRAAVTG
						CTSGKRLRLVDGTAISAPGGGS
						AEWRLHMGYDPHTCQFTDFEL
						TD\SRDGER\LARFAQPPDEIRIA
				-		DGEFVWRPKCIRSLAFGEADYI
!		1				VGVHWRGLRWLTAEGMRFDM
			•			MGFLRGLDCGKNGETTVMIGN
						SGNKKSP\GFPFRARFIAVSLPPE
						KALISKTRL\LSENRRKGRVVQ
						AETLEAAGHVLLLTSLPEDEYS
		1				AEQVADCYRLRWQIELAFKRL
						KSLLHLDALRAKEPELAKAWIF
		İ	Ì	İ		ANLLAAFLIDDII\SHRWISPPEV
						PDPKRRTNSLWRITKMVIWSLQ
		1				VAIRGTVSLTAPSLCNSACCCC
						VSTNSATTLNDFAFARSVIDLTI
25239	55607	Α	25377	2	333	RLGVWQDILSRRASCYHGMEV
						WDKDSFGHISFDDQSMGYSHL
						GHIVENSVIHYA\ARLVIGADGV
						DSWLRNKADIPLTFWDYQHHA
					ĺ	LVATIRTEEPHDAVARQVFHGE
						AFW
25240	55608	A	25378	3	235	VAVLEQRVHEPLAANAPPQLR
						VS/ARLGVWQDILSRRASCYHG
						MEVWDKDSFGHISFDDQSMGY
						SHLGHIVENSVIHYA
25241	55609	A	25379	1	774	
25242	55610	Α	25380	1	1076	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25243	55611	A	25381	1138	1737	SVSTEPSVAAFSSIKLRODPP/H
		į				DNLHHQYN*GDIAFC\FTPYFH
!		l				NCLCQS*WARAFQ*RANEF*R
		1				MNCTQLARTSGSHHLIPLTLRF*
						QICISSGQLIQYSVVCRFIHPPEH
						RLTIHPLAGQGVNLGFMDAAE
						LIAELKRLHRQGKDIGQYIYLR
	1	ŀ				RYERSRKHSAALMLAGMQGFR
						DLFSGTNPAKKLLRDIGLKLAD
25244	55612	В	25382	1	735	
25245	55613	В	25383	1	1941	
25246	55614	В	25384	847	849	
25247	55615	Α	25385	1	1227	
25248	55616	Α	25386	324	1533	EGFMEHQRKLFQQRGYSEDLL
						PKTQSQRTWKTFNYFTLWMGS
		l				VHNVPNYVMVGGFFILGLSTFS
		ł				IMLAIILSAFFIAAVMVLNGAAG
						SKYGVPFAMILRASYGVRGAL
						VPGLLRGGIAALMWFGLQRGS
					,	PFYDLIQTALSSPHKVTIEQFYR
						EVGVFLGIALIAVVISVLNNFFV
						SHYVFRWRTAMNEYYMANWQ
		ŀ		-		QLRHIKGRTAFINAIMTLIAFLP
					}	VLVTLSAHVPELPIIGHIPYGLVI
						AAIVWSLMGTGLLAVVGIKLP
						GLEFKNQRVEAAYRKELVYGE
		ŀ				DDATRATPPTVRELFSAVRKNY
					ł	FRLYFHYMYFNIARILYLQVDN
						VFGLFLLFPSIVAGTITLGLMTQ
						ITNVFGQVRGAFQYLINSWTTL
						VEL\MSIYKRLRSFEHELDGDKI
						QEVTHTLS
25249	55617	Α	25387	1526	1683	
25250	55618	Α	25388	2709	2918	CWCFWHRRSCCGSVRYALSAV
	1					PGCQCPET*RRRFPHSAF*RFRS
		l				SETGQTGAYVSGPDLQPERLQR
						LWC
25251	55619	Α	25389	1	1812	
25252	55620	A	25390	134	386	RLKPSAAIRRRVSSSRISTPLNPA
1			1			STTAFVGTDCA\IPRLRATSGLA
1					1	GSPLIQNRFCRARQPRWSAPVA
	<u>L</u> .					RHAAVYARHRWNQII

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25253	55621	A	25391	181	1630	RIRQFCFYHHCVLTLAICAFIVIT
23233	33021	^	23391	101	1030	LTIWVSFRVSVLVFAMADDIKV
						AVVGAMSGPIAQWGDMEFNG
1			•			ARQAIKDINAKGGIKGDKLVGV
						EYDDACDPKQAVAVANKIVND
						GIKYVIGHLCSSSTQPASDIYED
						EGILMISPGATNPELTQRGYQHI
		1				MRTAGLDSSQGPTAAKYILETV
						KPQRIAIIHDKQQYGEGLARSV
		1				QDGLKAANANGVFFDGITAGE
						KDFSALIARLKKENID\FVYYGG
						1
		ŀ				YYPEMGQMLRQARSVGLKTQF
						MGPEGVGNASLSNIAGDAA\EG
						MLAPVLITVDQPFVQAGFQPLR
İ						RLLPAHETGRRTRAYGVGRLT
						RQIRRQIATTFRHNAQAAESKD
						FQWYRCDPCQLFHLMNRQHAR
						QHHASNVKMAMIKVNRLFIGR
				1		RSLHRYVALNMRITLGGVLHH
1						GEVGEDQRVGTQLRRHIHGAL
						PTGVTVRMRKSVNRDVKFAA
		<u> </u>			50.5	MLMDKTHRFLQFFLGKVKAGE
175757	55622	IA	25392	2	1505	
25254	!		<u> </u>	 		
25255	55623	Α	25393	1	2038	GKOTSTSPAGWENSI RMRYT
	!		<u> </u>	 		GKQTSTSPAGWENSLRMRYT\ MSFSLIATGLI GVYLTTMPGY
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI AIADSDKLRVGRSGLNLEGLEN
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI AIADSDKLRVGRSGLNLEGLEN FIQTDASINRGNSGGALLNLNG
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI AIADSDKLRVGRSGLNLEGLEN FIQTDASINRGNSGGALLNLNG ENPIDQLKVVGRPHDRIDGPLK
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI AIADSDKLRVGRSGLNLEGLEN FIQTDASINRGNSGGALLNLNG ENPIDQLKVVGRPHDRIDGPLK TTGTARYAYEWHEEAPNAAYG
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI AIADSDKLRVGRSGLNLEGLEN FIQTDASINRGNSGGALLNLNG ENPIDQLKVVGRPHDRIDGPLK TTGTARYAYEWHEEAPNAAYG YIVGSAIAKGRLTALDTDAAQK
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI AIADSDKLRVGRSGLNLEGLEN FIQTDASINRGNSGGALLNLNG ENPIDQLKVVGRPHDRIDGPLK TTGTARYAYEWHEEAPNAAYG YIVGSAIAKGRLTALDTDAAQK APGVLAVITASNAGALDKGDK
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI AIADSDKLRVGRSGLNLEGLEN FIQTDASINRGNSGGALLNLNG ENPIDQLKVVGRPHDRIDGPLK TTGTARYAYEWHEEAPNAAYG YIVGSAIAKGRLTALDTDAAQK APGVLAVITASNAGALDKGDK NTARLLGGPTIEHYHQAIALVV
25255	55623	Α	25393	1	2038	MSFSLIATGLLGVYLTTMPGY WGILFVWALFGVTCDMMNWP VLLKSVSRLGNSEQQGRLFGFF ETGRGIVDTVVAFSALAVFTWF GSGLLGFKAGIWFYSLIVIAVGI IIFFVLNDKEEAPSVEVKKEDG ASKNTSMTSVLKDKTIWLIAFN VFFVYAVYCGLTFFIPFLKNIYL LPVALVGAYGIINQYCLKMIGG PIVLEKVLPPVVSVRVKGTASQ GQKIPEEFKQFFGDDLPDQPAQ PFEGLGSGVIINASKGYVLTNN HVINQAQKISIQLNDGREFDAK LIGSDDQSDIALLQIQNPSKLTQI AIADSDKLRVGRSGLNLEGLEN FIQTDASINRGNSGGALLNLNG ENPIDQLKVVGRPHDRIDGPLK TTGTARYAYEWHEEAPNAAYG YIVGSAIAKGRLTALDTDAAQK APGVLAVITASNAGALDKGDK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25257	55625	A	25395	356	849	GSACRVREIPRPASQMP/HTLPA GSRPPGSEAEPGPEAAPPPARRE AHQIPRPRRGHLPSSPGPRTHVA
						AHNIRVVTEAHADSHALLSDLG
						AAPFVVMSRILRVDGDEDPPHV
						DESIQPTTENAVVFRPVQLTSAI
		1				ETGGKEEPYGLSAASLGRAESS
		<u>.</u>				AASLAGRR
25258	55626	Α	25396	154	379	LVPRLKLGDLPSEINPLSSCSLV
						REKDPPTTSCTHT\TNPRNISPIS
		l				NPDSTGNRTVQLTW*PLPEPLE
						LWPKAL
25259	55627	Α	25397	4761	5127	LGSGDLPWEINPLSSCSLLREKD
						PPTTSGPQT\TSPRNISPISNPHTR
		1				TSKRLNRSGQAFLQNLLPQELA
			1			TSARNLTTRPRNACSPGFLLSR
		1				VPSVRDPTGNWTVQLTWQPLS
						EPLELWPKAL
25260	55628	Α	25398	1	753	MGSWLQGLRAMVFALNYMVT
		l				RVRNKKDPPTTSGPQT\TSPRNI
		ŀ				SPISNPR/PKETRFICGPKTPAPL
	Ì					MDWEGSLPLMFNHCRDTSLIIH
1						PCFQGVRPCRDACLSPSPLAASP
						AFLGKGQVPLNPFFTLSGKSRF
ŀ						SGGGASTPTPSFHVSTPSLLFW
						GRGKYPSTPSSPLVASPAFLGK
				1		GQVPLNPFFTLSGKSRFSGEGA
						KAPETITDAELRVTLTVEAAVC
						STIALSLGWEILPRHWGKEEVT
	}					KTIYYPVIP

SEQ ID		1	SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
1	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, (=possible nucleotide insertion)
		ļ		Jacque ii o		
25261	55629	Α	25399	174	2238	LRSGDLPWEINPLSSCSLLHEKD
•				1		PPTTSGPQT\TSPRNISPISNPRQR
						RHILSMDPKLRRRSWTWVGSLP
]			LVFNHCRDASLIIHPGFRGVRPR
						RDVCLGPSPLATITDAELYVTL
						TVEGKSIPFLIDTEATHSTLPSFK
						GPVALASIHVQNVQIKRDKEGH
						YIMVKGSIQQEELTILNMYAPN
						TGAPRFIKQVLNDLQRDLDSHT
						IIMGDFNTPLSTLDRSTRQKVN
						KDIQDLNSALHQADLIDIYRTL
						HTKSTECTFFSAPHRTYSKIDHI
						VGSKALLSKCKRTEIIINCLSDH
		l				SAIKLELRIKKLTQNHSTTWKL
						NNLLLNDYWVHKEMKAEIKM
		1				FFEINENKNTTYQNLWDTFKAV
						CRGKFIALNAYKRKQERSKIDT
						LTSQLEELEKQEQTRSKASRRE
						EITKIRAELKEIETQKTLQKINES
						RSWFFEKINKIDILLARLIKKKR
						EKNQIDAIKNDKGDIATDPTEIQ
						TTIREYYKHLCPNKLENLEEMD
						KFLDTYTLPRLNQEEDESLNRPI
						TGSEIVAIINSLPTKKSPGPDGFT
						AKFYQRYKEELVPFLQKYKEEL
						VPFLLKLFQSIEKEGILPNSFSEA
						SIILIPKPGRDTTKKENFRPISLM
						NIDAKILNKILANQIQQHIKKLI
						HHDQVGFIPGMQGWFNIRKSIN
·						VIQHINRTKDKNHMIISIDAEKA
		<u> </u>	2.5.100		0.6	FDKIQQPSC
25262	55630	Α	25400	3	267	EKDPLTTSGPQT\TSPRNISPISN
						LRLFAFTWTDPDTHQAQQITW
						AVLPQGFTDSPRYFSQAQISSLS
05062	55621	 	25401	167	400	VTYLSIILIKTHVLSLPIMSD
25263	55631	Α	25401	167	408	LGSRTFPWEINPPSSCSLLREKD
						PPTTSGPQT\TRPRNISPISNLVS
						GLFLLSSPTSLTIPQPLSSFNLDD
25264	55(22	 	25402	142	441	TLQSLPSLNF LRLGDLPWEINPLSSCSLLCEKD
25264	55632	Α	25402	142	441	PPTTSGP\PRNISPISNPVSGLFLL
					•	SSPTCLTIPQPLSPFNLGATLQSL
						PSLNFNSFHFLVETKETRFIRGP
25265	55622	<u> </u>	25402	70	201	KTPAP
25265	55633	Α	25403	79	381	LGSGDLPWEINPLSSCSLLREKE
						PPTTSGPQT\TRPRNISPISNPELA TLAGNLATGPRNARSPGFLLSH
						VLSVWDPTENQTVQLTWQPLP
						QPLELWPKAL
	L	<u> </u>	L		L	AI PEP MLKVP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25266	55634	A	25404	1	511	MWKGPKGLDMYGKSSVSPKTS
						DILGRDTLLLALKVQTVVLQTA
						CGEGHVAGNCGRPLETEGSLQ
						LTATKKLRDSVLQPKSPEFCQQ
						FTRAWNRTQVPDETEAPAGTY
						AAQSGDLPWEINPLSSCSLLHE
						KDPPTASGPQT\TSPRNISPILNR
						VSEVSDHAGTPALVLHP
25267	55635	Α	25405	570	1182	LGSGDLPWEINPLSSCSLLHEK
						DPPTTSGPQT\TSPRNTSPISNLH
						ALKGLKPAITRLLQHGLLKPINS
						PYNSPILPVLKPDKPYKLVQDL
						CLINQIVFPIHPVVPNPYTLLSSI
						PPSTIHYSVLDLKLAFFTIPLHAS
				1		SQPLSAFTWADPDTHQVQQIT
						WAVLPOSFTDSPHYFSQAQISS
						LSVTYLSIILIKTHVLSLPIVSD
25268	55636	A	25406	489	615	LRNTWNSPRLMACALHSGSPSC
,						TWGPYS*SGLSENLASQRVCPP
						YQNENVGRVRDTEIKEHCYEES
						SYRAATQIQVRKVGIITEPYPFM
						ASLRTHCIICETCYEMKISFSKS
					i	LNPLEKNKKSQPANCAVKEVS
					•	EQLRRRQAFSVAAGLKGKGSP
		ĺ				AQSIRKNLQKPYQHFQTLQQLL
						PLQWAPCYGPSNFATHGTHQG
						LWLVLSTVAAQAVPGALIAEA
						TALLAWIQGAVS
25269	55637	A	25407	579	715	LGSGDLPWEISPL\SCSLLHEKD
		l				PPMTSGPQT\TSPRNISPISNLR
25270	55638	A	25408	1553	1896	
25271	55639	Α	25409	964	1332	SCRFIRIYQGACCNLHPGEINSH
İ						VAHTKPVWWSLHTDTHENCHL
		1				NTAVFFYHWLALEFFLNPLRR/
						CSC*HSLPGQFPVSLCLRVLSVD
		1				CAHPWGKPEAPGS*HPGPWQFS
						SLPSPLWPLYQHP
25272	55640	В	25410	1	717	
25273	55641	A	25411	590	1142	PNSSWMRGEPKKDPPTTSGPQT
1						\TSPRNISPISNPR/PKETRFICGP
1						KTPAPLMDWEGSLPLMFNHCR
1		1				DTSLIIHPCFQGVRPCRDACLSP
						SPLAASPAFLGKGQVPLNPFFTL
		1				SGKSRFSGGGASTPTPSFHVSTP
		1				SLLFWGRGKYPSTPSSPLVASP
						AFLGKGQVPLNPFFTLSGKSRF
i	1	1				SGEGA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
į	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1		l		sequence		
25274	55642	A	25412	<u> </u>	790	MGLVHLQRQSYWTVQRSSINH
		-				FLGCNPHTGKDKTLYMRWTKS
	-	l				FRKVEDPPTTSGPQTDQPKKHL
						TKFKSETKETRFIHGPKTPAPV
	ļ					MDWEGSLPLVFNHCRDASLIH
						SHFKGVRPRRDACLGPSPLAAS
1						PAFLGKGQVPQPLISLCPDPLFP
						HPNLISLRPNPLCPHPDLVSLCP
						DPFPAFLEAHKNFQTPEPQQPGI
						PPEPPPPGACYKC/HEI*PPGQG
						MPAAQDSS*AMS/LSVRDPTGN
						QTVQLTWQPLPEPLELWPKAL
25275	55643	Α	25413	1	736	MOAMEKWKREAKVETEKDOK
23213	33043		25415		730	NQYQQTWLSDSASGYRYTKLP
						FYFRHNNLIPALAHLLFLDSLYS
		1				DLLTAAFFMQFWLSAQMSSDQ
				İ		RGLPGYLLWAEEISLLEISELTS
						GKQDSSWPEVKAGLKKEANYL
					<u> </u>	SSDSARLHPGEINSLVAHTKPV
						WWSLHMDTHEIWCRDSNWGD
						LPWEINPLSSCSLLHEKDPPTTS
						I .
İ						GPQT\TSPRNILPISNPPSSKHQE KPPEAISTLPNAAATAAAAVGS
į		ŀ				
25276	55644	A	25414	1644	2010	LLWSL LGSGDLPWEINPSSSCSLLREKD
23276	33044	A	23414	1044	2010	PPTTSGPQT\TSPRNISPISNPHTR
						TSKRLNRSGQAFLQNLLPQELA
					ļ	TSARNLTTRPRNACSPGFLLSR
						VPSVRDPTGNWTVQLTWQPLS
						EPLELWPKAL
25277	55645	A	25415	1	1035	MGKEYSLDVKKSIKVFKDNRS
23211	33043	^	23413	1	1033	QGNRLQKLGLEDTDREDAMGF
İ		1				GSHRAKLTVVAALGACHCPEN
		1				EGQTVLPSSTSGRSDKRERVSA
·					<u> </u>	GYTPPFFVKEGVPSPLFKLHVV
						RCCSIQSHHSRRSQNCAEDVIR
					1	KTKTDQAVSSMSSAPVTVQQL
		i				NIIPALLVALTCGWAFCLMVSQ
						`
						EQYDEASKQAASFSNNAVIPVY
						MSHMSKTLYKSEVFPINTYAQK
						ICFIAHTKPVWWSLHMDAHEI
						WCRGRRSTDLPWE\INPLSSCSL
		1				LREKDPPMTSGPQT\TSPRNISPI
		1				LNQELATRAGNLATRPRNACSL
		1				GFLLSRISSVWDPT*KFGLVQLT
		1		<u> </u>		LGKPLPEPLELWPKAL
25278	55646	Α	25416	174	393	LRSGDLPWEINPLSSCSLLHEKD
		1				PPTTSGPQT\TSPRNISPISNPGET
						KETRFIRGPKTPAPVMDLGRQP
				1		SLGV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l '	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25279	55647	I .	25417	2	4574	WDVLGAVIGTECGTIESGLSMV
23219	133047	Α	23417	2	4374	
İ	-	ŀ				FLKDGERKLCTPSMDTTGYGN
					<u> </u>	LRFYFVMAKWIYIPENVVKGR
		İ				KEIILVLDFTWCMVHRGICDPG
						NSHENDIILYAKIEGRKEHITLD
						TLSYSS/L/SSLYWPVMSQILFW
						QRTVH*TWL\QL*DGIPDIPNVY
						F*KLWQFQALLLP*LLLYPWC*
						SQLWLWCLGQW*GPGFQQRW
						AASANYIFP*QLTIQNNLRRTTR
						*CKAVWNSVQMVATVSFFPER
						RCMGY**DYHDICA
25280	55648	Α	25418	1	363	
25281	55649	A	25419	1	301	
25282	55650	A	25420	2	309	
25283	55651	A	25421	3	512	YIGPSCLKFCSGRGQCTRHGCN\
						CEMASQTFPMFISESFGSSRLSS
						YHNFYSIRGAEVSFGCGVLASG
						KALVFNKDGRRQLITSFLDSSQ
						SRIISVELPGDAKQFGIQFRWW
						QPYHSSQREDVWAIDEIIMTSV
						LFNSISLDFTNLVEVTQSLGFYL
		ŀ				GNVQPYCGHDWTL
25284	55652	Α	25422	2	878	<u> </u>
25285	55653	A	25423	1	750	
25286	55654	A	25424	1	1404	

SEQ ID	SEQ ID NO:	L	SEQ ID NO:		i -	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	•		,	sequence		
25287	55655	Α	25425	1	2506	GCSRLLVTVDLNLTNAEFIQFY
						FMYGCLITPNNRNQGVLLEYSV
						NGGITWNLLMEIFYDQYSKPGF
						VNILLPPDAKEIATRFRWWQPR
						HDGLDQNDWAIDNVLISGSAD
						QRTVMLDTFSSAPVPQHERSPA
1		ŀ				DAGPVGRIAFDMFMEDKTSVN
1						EHWLFHDDCTVERFCDSPDGV
İ						MLCGSHDGREVYAVTHDLTPT
					1	EGWIMQFKISVGCKVSEKIAQN
			ļ			QIHVQYSTDFGVSWNYLVPQC
			ŧ			LPADPKCSGSVSQPSVFFPTKG
						WKRITYPLPESLVGNPVRFRFY
						QKYSDMQWAIDNFYLGPGCLD
						NCRGHGDCLREQCICDPGYSGP
						NCYLTHTLKTFLKERFDSEEIKP
						DLWMSLEGGSTCTECGILAEDT
						ALYFGGSTVRQAVTQDLDLRG
1						AKHDYILLPEDALTNTTRLRW
						WQPFVISNGIVVSGVERAQWA
						LDNILIGGAEINPSQLVDTFDDE
						GTSHEENWSFYPNAVRTAGFC
						GNPSFHLYWPNKKKDKTHNAL
						SSRELIIQPGYMMQFKPFGFCRS
	-					DSWQLVQTQCLPSSSNSIGCSPF
						QFHEATIYNSVNSSSWKRITIQL
						PDHVSSSATQFRWIQKGEETEK
						QSW\AIDHVYIGEACPKLCSGH
						GYCTTGAICICDESFQGDDCSV
						FSHDLPSYIKDNFESARVTEAN
						WETIQGGVIGSGCGQLARYAH
						GDSLYFNGCQIKRAATKPL\DL
25288	55656	Α	25426	146	10561	GGGGGGGGGMERSGWARQT
İ						FLLALLLGATLRARAAAGYYP
						RFSPFFFLCTHHGELEGDGEOG
		İ				EVLISLHIAGNPTYYVPGQEYH
						VTISTSTFFDGLLVTGLYTSTSV
						QASQSIGGSSAFGFGIMSDHQF
						GNQFMCSVVASHVSHLPTTNLS
						FIWIAPPAGTGCVNFMATATHR
						GQVIFKDALAQQLCEQGAPTD
						VTVHPHLAEIHSDSIILRDDFDS
						YHQLQLNPNIWVECNNCETGE
						QCGAIMHGNAVTFCEP
25289	55657	A	25427	1	296	QCG/HMHOHWA IT CEI
23209	123037		23721	l •	270	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25290	55658	A	25428	1	557	MARSLGVLVALPFPLPVGFNAE
						EAHNIVKEFSPGKSHPRESLSQQ
						RETLVQAGAFHWFGWRGKEK
:						LKPKFNTFGRLPNEMLEQWLSP
						TNQCVDGV/LKGGED\YNHNNI
						NQWTASIVEQSL\THLVKLGKA
						YKYIVTCAVVQKSAYGFHTASS
						CFWDTTSDGTCTVRL/WENRT
						MN\CIVNVFCHCLFVL
25291	55659	Α	25429	178	2270	VRRLFCVQTGGSPVGRGSLHHL
						WLVFSHVLSLLCSPCLLAPGM
						WGFHDRDLVLRKALYTMMRT
						GAEREALKRRWRWQQTQQNK
						ESGLVYTEEEWEREWTELLKL
						ASSEPRTHFSKNGGTGGGHLPP
		ŀ				EGKPLHGCSSLLRSGKGGHVK
						AEPSRLQQSQAKPPLTSQIRALH
						FPIPSQINEKPGAGHWSAKLHFP
						YWQLGEAGEEGSSPPDIMRCL
						LTCLSVPSTAVIPLTDSEHKLLP
						LHFAVDPGKDWEWGKDDNDN
						ARLAHLILSLEAKLNLLHSYMN
		ļ.				VTWIRIPSETRAPLAQPESPTAS
						AGEDVQSLADSLDSDRDSVCS
						NSNSNNGKNGKDKEKEKQRKE
						KDKTRADSVANKLGSFSKTLGI
						KLKKNMGGLGGLVHGKMGRA
						NSANGKNGDSAERGKEKKAKS
						RKGSKEESGASASTSPSEKTTPS
						PTDKAAGASPAEKGGGPRGDA
						WKYSTDVKLSLNILRAAMQGE
						RKFIFAGLLLTSHRHQFHEEMIG
						YYLTSAQERFSAEQEQRRRDA
						ATAAAAAA\PPPPPRP/ERPPRRP
						ETEGVPVPERASPGPPTQLVLK
						LKERPSPGPAAGRAARAAGG
1						TA/CPGGRRAACERQRTSAWPQ
						PPGASAPERHPRAGVGRAGRG
						VRAGRGGAAAVRHVPAAEPLA
			1			VVAELQPGARRRPAHRQHGRV
		<u> </u>				AGARGARGPTGRGGDRGGGRA

SEQ ID			SEQ ID NO:		•	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1 '	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25292	55660	Α	25430	404	1116	SLGPLMLSGQKLQTSYTMEYY
						AAIKR/DKIVSLVGTWLELEAIIL
						KKLMQEQKTKYHMFLLDSLSH
						GMQQAERVLIADGTMLLSAQY
						QFENNLLVPILRVSFRKSLQRKE
						EPGRKAKKSQTSGEGIKKRRFW
		l				FEWKEPVALSSQGWYEAKGSE
						RPEKGHLGLLRAESFQDLRTAL
						ISKPYFSVPGKVGPAIKCPETLP
		l				ELKTYPQYGVTLGEPVRNEQQ
						QASSFPSLSNTHLSKKRNES
25293	55661	Α	25432	1571	1782	WLLCRLPPRVPDQL*WCGCQPE
		1				KCAGAEFQDGSAMPLPSRTCPH
						SPRMCTGWSTTCATTDSATTA
		ŀ				CPLAL
25294	55662	В	25433	69	1160	
25295	55663	Α	25434	14	437	LQTSHPYPTTSKC/PPP/CEKPSP
						TSL/C*RRSAPHLQPPE
25296	55664	Α	25435	782	997	MMLCQLQRPV/CGSHSCSRG/C
						PPAQ/CPSPVPVPSPQSAFPVPQS
				ŀ		PSSTCPNPSHPIHPIPMVPSSWSG
						HWDSK
25297	55665	A	25436	1117	1815	GDAHTEQCLQNLYPRGSRPPLP
						LESHPSSSKHSPKACPTGRS/CG
						PWK/CPGPEVRP*PPPSCTR*DP
						HFRGLLEGLLEPRHVTCLALCG
						PQRVLYWPVYHERLGEGWDW
				1		CPSSWPGA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25298	55666	Α	25437	1261	2757	ALGKDEKSQGPPIQTGGLGPRL
						FRLPSSQGQERRFAAAKAPSSL
						VPHNGSPLSCWRGLREEGGLSL
						ATLWTPCGVGSGVPAAPPSAW
						CDLGKLLVCWRGQNRGPRSTQ
						TDVRRHEARSRQRRRKCVAAP
İ						VAPRSRRWRCPAALAAPAVAV
						PVVFALIFLLGTVGNGLVLAVL
	1					LQPGPSAWQEPGSTTDLFILNL
	1					AVADLCFILCCVPFQATIYTLD
						AWLFGALVCKAVHLLIYLTMY
						ASSFTLAAVSVDRCAVPG\HPL
						GRAVGAGVGGGVLNVGLVWL
						LAALFSAPYLSYYGTVRYGALE
						LCVPAWEDARRRALDVATFAA
						GYLLPVAVVSLAYGRTLRFLW
						AAVGPAGAAAAEARRRATGRA
						GRAMLAVAALYALCWGPHHA
						LILCFWYGRFAFSPATYACRLA
						SHCLAYANSCLNPLVYALASR
						HFRARFRRLWPCGRRRRHRAR
						RALRRVRPASSGPPGCPGDARP
						SGRLLAGGGQGPEPREGPVHG
						GEAARGPE
25299	55667	A	25438	1	1624	MKDKSKTLGLEASIFRNNWKC
				-		KSIFEGLKGHQEGYFSQMIISYE
-						KIPSYRKSKSLTPHQRIHNTEKS
						YVCKECGKACSHGSKLVQHER
	İ					THTAEKHFECKECGKNYLSAY
						QLNVHQRFHTGEKPYECKECG
						KTFSWGSSLVKHERIHTGEKPY
						ECKECGKAFSRGYHLTQHQKIH
İ		ŀ				TGVKSYKCKECGKAFFWGSSL
						AKHEIIHTGEKPYKCKECGKAF
			1			SRGYQLTQHQKIHTGKKPYEC
}		1				KICGKAFCWGYQLTRHQIFHTG
		İ				EKPYECKECGKAFNCGSSLIQH
-		l				ERIHTGEKPYECKECGKAFSRG
						YHLSQHQKIHTGEKPFECKECG
						KAFSWGSSLVKHERVHTGEKS
						HECKECGKTFCSGYQLTRHQV
1						FHTGEKPYECKECGKAFNCGSS
						LVQHERIHTGEKPYE\CKECGK
					1	GFS\GGYHLTQHQKIHTGEKPF
					1	KCKE\CGKAFSWGSSLVKHERV
		1				HTNEKSYECKDCGKAFGSGYQ
1						LSVHQRFHTGEKLYQRKEFGK
						TFTHGSKLVHERTHSNDKPYK
						YNECGEAFLWTTYSNEKIDTDE
	ــــــــــــــــــــــــــــــــــــــ		L		<u> </u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	ł .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25300	55668	Α	25439	32	447	VLGGSMMFREGQRGNRQKGP
	ŀ			1		KKRRNLALSSALLASTA\PAGG
						QPRHWRPG*CCLRQQL*PLQGN
						WAPVSPLQGHRSQLRAGYSISS
		ł				FLQMFAP*VQGAARAQPAPGA
						AVPRGSGRQGGPRAVDLHGFR
		1				VGWNRLRAASM
25301	55669	Α	25440	1	858	
25302	55670	Α	25441	147	1088	SRTTYKGKSSFQTYSDYLRWES
						FLQQQLQALPEGSVLRRGFQTC
						EHWKQIFMEIVGVQSALCGLVL
						SLLICVAAVAVFTTHILLLLPVL
						LSILGIVCLVVTIMYWSGWEMG
		1				AVEAISLSILVGSSVDYCVHLVE
					:	GYLLAGENLPPHQAEDARTQR
:						QWRTLEAVRHVGVAIVSSALT
		1				TVIATVPLFFCIIAPFAKFGKIVA
		İ				LNTGVSILYTLTVSTALLGIMAP
						SSFTRTRTS/CPQGPGCRAAGRG
		1				PGAGCLPRAPAERL*DSPARRG
						LPIARDGLWTLAPLVPWVGDR
						SCFPARLQLAVSPGLGPGRPAG
25303	55671	Α	25442	1	1596	MGVSVRVDCGCARASEFLGRP
		1				RGAVRGADPGSGLTETCAQSR
j		1				TWARARSKGGFRLTLNFYGPR
		l				YPSEGTSRGQRSEALPFLDIQHP
		1				TPLGRPRMLQSRGEMVRSWGP
		1				VVEGKLGFCPPLTVFPGGLRGP
		1				HPGLAGERENFARGAKQEACL
		1				LQGPGDAGLTHPCPATGYFPEA
		l				SCTSRSIRMFINSQERQLADMSL
						FIADSWLCYSGIICWKSLEHQR
						GASLSCVYMEGKGGNIWMKAS
						NVLPLCQMLPTSGHSKSDLFGG
		ļ				LAVEKQMRQDHHILLHDRSGF
						QLAVTSPLLTRLGKGHIIQVPV
						YVLDAPSRWLGGLRLFRDSSCG
						LVLSLLICVAAVAVFTTHILLLL
	1					PVLLSILGIVCLVVTIMYWSGW
]						EMGAVEAISLSILVGSSVDYCV
						HLVEGYLLAGENLPPHQAEDA
						RTQRQWRTLEAVRHVGVAIVS
						SALTTVIATVPLFFCIIAPFAKFG
						KIVALNTGVSILYTLTVSTALLG
						IMAPSSFTRTRTS/CPQGPGCRA
						AGRGPGAGCLPRAPAERL*DSP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25304	55672	A	25443		2355	MDHPGFREFCWKPHEVLKDLP LGSYSYCSPPSSLMTYFFPTERG GKIYYDGMGQDLADIRGSLEL AMTHPEFYWYVDEGLSADNLK SSLLRSEILFGAPLPNYYSVDDR WEEQRAKFQSFVVTYVAMLAK QSTSKVQVLYGGTDLFDYEVR RTFNNDMLLAFISSSCIAALVYI LTSCSVFLSFFGIASIGLSCLVAL FLYHVVFGIQYLGILNGVAAFV IVGIGVDDVFVFINTYRQATHL EDPQLRMIHTVQTAGKATFFTS LTTAAAYAANVFSQIPAVHDFG LFMSLIVSCCWLAVLVTMPAA LGLWSLYLAPLESSCQTSCHQN CSRKTSLHFPGDVFAAPEQVGG SPAQGPIPYLDDDIPLLEVEEEP VSLELGDVSLVSVSPEGLQPAS NTGSRGHLIVQLQELLHHWVL WSAVKSRWVIVGLFVSILILSLV FASRLR\SASRAPLLFRPDTNIQ VLLDLKYNLSAEGISCITCSGLF QEKPHSLQNNIRTSLEKKRRGS GVPWASRPEATLQDFPGTVYIS KVKSQGHPAVYRLSLNASLPAP WQAVSPGDGEVPSFQVYRAPF
						GNFTKKLTACMSTVGLLQAAS PSRKWMLTTLACDAKRGWKF DFSFYVATKEQQHTRKLYFAQS HKPPFHGRVCMAPPGCLLSSSP DGPTKGFFFVPSEKVPKARLSA TFGFNPCVNTGCGKPAVRPLVD

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25305	55673	A	25444		1737	MLYAAECLRNVMKLQVIPSAL NEALQAQSTWSDLGLASHALS STWCLHGESAAWSQPAVVECG QPRDQLQTHSEAWLCALVATF SQELEQFAVTGAQISDRQGHAV CTAPPRGRPSTLMTNLVCASSA DECGGGKTETQTAGGRLQGAE TRSRGQLTRLITRTQAAEPRWF LQLLKRSANRSRALTRPFSGLSS DGARRLRLSFVSAVSSADTKPT VPVSSDSSGRDGCGETRAEAGP GGPHVAQPQRVAPPAAAARDF RKPRGLNRQNVSP/PGPGEQGP G/PRPQKPAGRAP\PEPQGEPGL GLFPR\GRPPRSFPGSPADPAAS VSHTQGPSPDTGLPGTCQNRLP NPAAVDEQTCTRGSLNCRLSRS PRVSERRGPEGTPGRRPAAPPQ CAGTRGLSPRADLGAAIRTPRA PYRPHGQGRPSRHNCDSVSSQG WPPHSCWWKTTSNGMPWKAS FCSRTAQPANQNTGNHERGLRS VAMFGDRASREATPALSLCESQ GCFEDEKEVFSPEPHVETSDEPL
						PICSGPGDGWGDPIARPGRGNN LSPGTSAWEVPRWIRVTGEDW LSTGPAECVFINDLTT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25306	55674	Α	25445	2	1642	LLSSISAWFGPPASTPAFTMSIR
		1				VTQKSYKVS\TSCPRAFSTRSYT
		Į	į			SRPGSRISSSSFSRVGSSNFRGDL
						GGGYGGASG\MGGITAVTVNQ
		1				SLLSPLVPEVDPNIQAVRTQEKE
						QIKTLNNKFASFI\DEVGFLEQQ
						NKMLGTKWSLLDQQKTA\RSN
						MDNMFESYINNLRRQLETLGQ
		1				EKLKLEAELGNMQPLVEDFKN
						KYEDEINKRTEMENEFVLIKKD
						ADEASMNKVELGSRLEGLTDEI
						NFLMQLYEEEIRELQSQISDTSV
						VLSMDKSRSLDMDSIIAEFKAQ
						YEDIANRSRAEAESMFQIMYEE
						LQSLAGKHGDDLRRTKTEISEM
						NRNISR\LQAEIEGLKGQR\ASL
		1				GGRPLQDAE\QRGK\$WPIK\DA
						NAK\LSEAGSPPLQAGPKQDMG
						R\QLREY\QKLMNVKAGPWTIE\
						IATYQESCLEGEEEPGWSLGCQ
						NMSI\HT\KTTQRGYAGRPEPPP
	-	l				MGGFTSP\GFSYKPGAPGFGSG
						AGS\SSFNRHQLSPGAVVVKKIE
						TRDGKLVSES\SNVLPKVKQLR
		Ì				QPLPSLPLLRCPRGLGKEAPYP
25307	55675	Α	25446	1	1280	MAYLCNAICMTHMVTLDPVDQ
		l				ALPGATTATSVPGPQCDASCM
		ŀ				MLFAVVPFNGIALAATQGAQS
	İ	l				MLPPRVQGSLSSKPSTRRGRDA
		ļ		İ		PGLQVSQAPPPRRGFAVGRRYS
		l				PPALAPGRCAAPHGGGRKELPT
				i.		RRPGHGMAPKFPDSVEELRAA
						GSESFRNGQYAEASALYGRAL
						RVLQAQGSSDPEEESVLYSNRA
						ACHLKDGNCRDCITDCTSALAL
	1	l				GPFSIKPLLRRASAYEALEKYP
	1	l				MAYVDYKTVLQIDDNVTSAVE
	İ					GINRGPGLKEDTTQKRWNSLPS
						ENHKEMAKSKSKETTATKNRV
			ł			PSAGDVEKARVLKEEGNELVK
						KGNHKKAIEKYSESLLCSNLES
						ATYSNRALCYLVLKQYTEAVK
						DCTEALKLDGKNVKAFYRRAQ
						AHKALKDYKSSFADISNLLQIEP
						RNGPAQKLRQEVKQNLH
			L	1	L	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25308	55676	A	25447	1	1297	MHLLSAYFVPGSMLVGSGYM MVNITDVSPPSWSFLCSMLITTT
						QGAQSMLPPRVQGSLSSKPSTR
	ļ					RGRDAPGLQVSQAPPPRRGFAV
						GRRYSPPALAPGRCAAPHGGG
						RKELPTRRPGHGMAPKFPDSVE
		İ				ELRAAGNESFRNGQYAEASAL
						YGCALRVLQAQGSSDPEEESVL
						YSNRAACHLKDGNCRDCIKDC
						TSALALVPFSIKPLLRRASAYEA
						LEKYPMAYVDYKTVLQIDDNV
						TSAVEGINRMTRALMDSL\GPE
						W/RALKLPSFPLVPVSAQKRWN
	}					FLPSENHKEMAKSKSKETTATK
						NRVPSAGDVEKARVLKEEGNE
		1				LVKKGNHKIAIEKYSESLLCSN
		1				LESATYSNRALCYLVLKQYTEA
		1				VKDCTEALKLDGKNVKAFYRR
						AQAHKALKDYKSSFADI\SNLL
		<u> </u>				QIEPR\NGPAQEVAGREVKQNL
25309	55677	Α	25448	3	491	GITFTHSPAATGGGAQRVRRRD
						PPPGVQLRNMAQETNHSQVPM
						LCSTGCGFYGNPRTNGMCSVC
8						YKEHLQRQNSSNGRISPPATSV
						SSLSESLPVQCTDGSVPEAQSAL
			ŀ			DSTSSSMQPSPVSNQSLLSESV\ ASSQLGQYICGTKAVPETEDVQ
		1				ASVSDTAHV
25310	55678	A	25449	1	860	HLLLLAIDYNLPIAGVLMESQY
23310	33076		2377)	1	1000	LGASPKRLVIPNLYAHRKSLAT
						LEPYAPRRPMPIARRYAIPYMS
						HHNGYIGDVYYRLTVEEKKSSF
						IARRSILTTEFTIRRTKKVPVPSI
						NMSSASPLEDESRPSAGVQLRK
ļ						GRYRLLRDRYRPGCNSVGESPR
						KSSFSDDIMATESRESIFAEEPR
:						TGEHSVRGPVSNQSLLSESVAS
						SQLDSTSVDKAVPETEDVQDSL
						NASPTKQQCG/DVQTVDLYSNN
		İ				KELFAQKCYRHRYISRLKKCCL
						CHELVPRDTIKSPESLSVGS
25311	55679	Α	25450	880	1196	SPTTRRGTPGTCLKPTACAGAA
						*PGCSARRTCASAAPLSTCPPSS
		1				CAAPPPAPAAVPSTPRPTSPSPW
						AAPASPSRRRTQTASTPASTNR
						APSSCWAPTTRPLAP
25312	55680	Α	25451	254	579	IEKISLEPKNRPSPQIPNLVGP/R
						LFLVPGDFQSQVPNFLPFKPSSG
						PHPAAGKGQPFSFLPAWRSTGP
						ERATPGPGRPAACSALEAPSLR
	<u> </u>		<u> </u>			GPSGYSAQEPPRRPISARI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
ļ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
25313	55681	A	25452	6	1442	RSELIRHSWPRHAQLRGARART
						AVSIRECISVHVGQAGVQIGNA
		İ				CWELFCLEHGIQADGTFDAQAS
				}		KINDDDSFTTFFSETGNGKHVP
						RAAMIDLEPTVVDEVRAGTYR
		İ				QLFHPEQLITGKEDAANNYAPG
		1				PPTPGGKESIDLV\LDRIRKLTD
		1				ACSGLQGFLIFHSFGGGTGSGFT
						SLLMERLSLDYGKKSKLEFAIY
						PAPQVSTAVVEPYNSILTTHTTL
						EHSDCAFMVDNEAIYDICRRNL
						DIERPTYTNLNRLISQIVSSITAS
						LRFDGALNVDLTEFQTNLVPYP
		1				RIHFPLVTYAPIISAEKAYHEQL
						SVAEITSSCFEPNSQMVKCDPR
		1				HGKYMA\CCMLYRGDVVPKD
						VNVAIAAIKTKRTIQFVDWCPT
		ŀ				GF\KVGINYQPPTVVPGGDLAP
1		:				VQRAVCMLSNTTAIAEAWARL\
						DHKFDLMYAKRAF\VHWYVGR
						GDWKKGEFSEAREDLAALEKD
İ					ļ	<u>}</u>
25214	55692	 	25452		618	YEEVGVDSVEAEAEEGEEY SSAGREPDPSDLPRRLCFTHRLP
25314	55682	A	25453	3	018	AARRWVQLCVHASPEPGGQGV
						CPGRSERMVIRVFIASSSGFVAI
						1
						KKKQQDVVRFLEANKIEFEEVD ITMSEEQRQWMYKNVPPDKKP
1						TQGNPLPPQIF\NGDRYCGLLVS
1						FSVICIKNPNWHNLFLLIPWSGI
1						NPYCEQDSSFSENVGDEILKNS
						GVPRDFTSILIQKERKSEKVKG
25015	55600	١.	05454	2	404	VDQGHR
25315	55683	A	25454	3	424	SWTMAATIQAMERKIESQAAH
						LLSLEGQTGMA*S*GTPPCCPG
						AAAGSRSSCCAAGSPRSPAASA
						AARTPAAGSPARPTCPPAGSRT
						PRLSSRNQPASSRPSRFDLLGKA
						GEQPAIQSSSPWPESWLPSSTK
25316	55684	A	25456	268	445	
25317	55685	A	25457	3	205	
25318	55686	A	25458	27	272	AKCSGQGPGRVRAQGERADHE
23310		[*	23 .50	 -'	[-,-	CPGAGAGGGGCNSAASAREES
						SRLISG\SLQMQNVSLASRRKEE
						EVRAAAAGGAAAGRPRV
25319	55687	A	25459	1	253	2.70mmioommioom
25320	55688	A	25460	2145	2399	
23320	133000	1'1	123700	14177	1-3//	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25321	55689	A	25461	174	1309	HLLCCRAORRPOTPPAARGLEA
						CSEGALKMPEAPP\LLLAAVLL
						GLVLLVVLLLLLRHWGWGLCL
						IGWNEFILQPIHNLLMGDTKEQ
						RI\LNHVLAACGSPGNAQSVLG
						GHLTPNCEQKEW/APLNVGDK
						KGQDSWDALISREQPALPCLLG
						NLGAYFGLLQLLRMAR\LLVNQ
						GARLIHHSRFNPDLCPPSPKRM\
ŀ		ŀ				VDFAGVKDKALFQVTL\VCGAS
						PETFHPPSLKKKYDVGHNWTW
						VFLDHWKDRYL\PDTLLLE\EC
			ŀ			G\LLPEGGQLLLAEQRLSSPGAP
						\DFL\AHVSREPACFG\CT\HYQI
1						VPWKYRE\VVDGLGRRPI\YKG
						P\GSEAGALTAPPRPPSRALSPK
						PGTEGARPWLLLTLVRLRACVL
						NAKSHARAESLRA
25322	55690	A	25462	2470	3460	SGPCPRCCAECRPYKTRPRSRV
						PGTHGPGPADHGQPPCCMAGA
ŀ						HPPRPQAWMLLQTHSQDCEGK
}						VGCGGIFCPCYHHCKHTHHHH
		1				ННРИНННҮНҮНННИННОННН
						ннниннуннунзнуннуннн
						НСННН\НННННРҮРННҮННН
						RGQDHНРНРНННН/НННҮННН
						НҮНННҮНРНННННОРОН
		1				НННОНУНННУНННРННРНН
1]		РНННННҮНҮННННННҮ\НРҮ
						РННҮНННННННРНРНННН/
	8					НННҮННРННПҮННННРН
						НННКРНННКНОНРННҮННН
						НРНРСНННКІНННКНННҮНН
						TSSVQGPGSQPAGASLPGQACR
25323	55691	Α	25463	1	407	DAERQEALGIVRRIGTDTEAAT
						EPAGATVPAAAAAARIGTVGP
						QPPAMPRRKRNAGSSSDGTEDS
						DFSTDLEHTDSSESDGTSRRSA
						RVTRSSARLSQSSSRISRSCSKS
		<u> </u>			<u> </u>	GSLLALRSLLTLPEE*PVVSSSL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25324	55692	Α	25464	3	2539	RILYIPDGHRAPGASQSGNWEH
ļ.						PLLLLEPGNLASSPSMSLAYPHP
						CGLTIHQEHRRAHEELAYTPND
		l				RASSTGHPAFIEDGNPSPVLAFA
ŀ	ł	İ				ASPRPNDSYILKREPPESCEKER
						VFEEATTRGKYGEGAKQETFTF
						ALTLVFIQCVINAVFAKILIQFF
						DTARVDRTRSWLYAACSISYLG
						AMVSSNSALQFVNYPTQVLGK
						SCKPIPVMLLGVTLLKKKYPLA
						KYLCVLLIVAGVALFMYKPKK
						VVGIEEHTVGYGELLLLLSLTL
						DGLTGVSQDHMRAHYQTGSN
						HMMLNINLWSTLLLGMAVSCP
						DQGPELVPRCPFVQALEKPSWK
						NLHQQDLFATCYWHNEGESCV
						SCHGKTLSSKTQGGILFTGELW
						EFLSFAERYPAIIYNILLFGLTSA
						LGQSFIFMTVVYFGPLTCSIITTT
						RKFFTILASVILFANPISPMQWV
						GTVLVFLGSTQIILDNLKILNTS
						AKTLLYKTPLSWQELEGERASS
						CAHKRSASWGSTDHRKEISKLK
						QQLQRTKLSRSGKEKERGSPLL
						GDH/VSAGSTEGVPS/RASPQGP
]	ŀ				LSCDSAPACT/VSLEGLNQELEE
		İ				VFVKEQGEEELLRILDIPDGHR
						APAPPQSGSCDHPLLLLEPGNL
						ASSPSMSLASPQPCGLASHEEH
						RGAAEELASTPNDKASSPGHPA
						FLEDGSPSPVLAFAASPRPNHSY
]						IFKREPPEGCEKVRVFEEATHQ
25325	55693	A	25465	845	1316	LSLGDSAQCLLPHASWCQVAG
						HPAFLEDGSPSPVLAFAASPRPN
						HSYIFKREPPEGCEKVRVFEEAT
						S/RRVLTGPFLTSCPDK\NKVHF
						QP\TGSAFCPVNLMKPLFPGMG
		}				FIFRNCPSNPGISSSPGQPQATTS
ĺ						EGSGSLQGLPTAIRAMAAHPYH
		•				Q

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25326	55694	A	25466	1077	1506	CDACGRRCRTCRTFCLGLQPAG LDGHHHRHPDCWFPWSPRHR* GMIIDVFSRYSGSEGSTQTLTKG ELKVLMEKELPGFLQILTNEDE PLVTIPFLSHSCSLAYLLTTSGN RVNALLKWLLKLGERWQMTE EGIKGQRSGHGPEQAENPLGDS VPQDTTKDVPLTQHRLHQGLE AGVPKTSVPARPGLPVPALQHA QIPGPCERALLHCHNGPQRLQH SLQGEEQRETHLRGQDWPCTL QMRTLKREVHALPGGRAGGAA REGLGASICETGEHQMTNFQVP RWPPCFPGQAGPVLGTGTRHG LDLLSGRPGTVRKGSSVARIMK EEVTVTKQSAADLRHTGQGPT CCVLSGGRCGSQACSRRPFSLL LGDESEPHGRGPFAAALHHRR NPLNDVMLVGDDAGPAGHFAS VYNPLAWTVTTIVTLTVGFPGV RVTDEAGHPVPSQVSGIQRCLQ GMKAPPVTVTVSGEKGSCLCA SCVLCLEHPYFPLVPWLPPVFR DSVIPSSTSPQEAVLADVGFRN NVPQRAQLKPQKQILTPF
25327	55695 55696	A	25467 25468	846	783 1392	PDPAAHRFGHPTLHPGSLRTSS GAGTWYQNPQHHNDHDYYHH ННННОДННҮНҮННННННННН ННҮННННННННННННННН
25329	55697	A	25469	181	412	LRSDQQTRKFVSRSSGACSPPTT TELSWPPSLLPSL*FHSPSSQP* MNHSASTTSPLPSFITERESILAT EWDTARN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25330	55698	A	25470	1	848	MPSLNTIVSPLGSPHTSLIPPML
						LSANLPFDTTIMIKLQTACSAPI
						HNLQRPCGENLRTSTFVRRLDR
1						NLVKSPQPGGGGQPSRQWEAR
						GAEPQGRRDPSEVQ/RPAGRGF
						GRISRAPRKPSRAPTPTQGLGPP
						LGAGGQGSLASCRPARKNARPP
						TRAAAHSAHPHSHWHSHARPQ
						PQSPSAAGQSGDPWPPARVPPP
						PLLRLPQRRTAGRTPVDAPARG
						IFTLSLSSSHSKSSGKHRGARKT
						PEMWPLFRGSLSKLSRRRKVSP
		1.				SAALHWRLRRTQKRLCC
25331	55699	A	25471	92	1083	TMRTHGHRKGNITPWGLSGVG
						GL/ETGRALGEIPNVDEGLMGVI
						AAVAIVLVSLLFVMLRYMYRH
						KGTYHTNEAKGTEFAESADAA LQQRERALDSSRDSHLETLGAC
						PGRNGGGLALQEATPRDPGRR
						WPPQRPPFAPRRWEKIWAHGPL
						GSAGQHQLTGRKVLVEGLGCI
						CMAHTDLQNTSVAKMAINRPN
						EYAQVSLENDVSCKVKAVQAP
						QRLHWPSVDADTGGRGAVHIA
						FPKHVEPETPFAHTIPARNSWT
						KWSSEHILETLSPQKPALPWAP
					1	RQQGRPLGLTAPISKRLGPGDG
						PEGEAGERLAQAHFINTRPPST
						GPPGTPAAD
25332	55700	Α	25472	73	424	RPGMWSTRSPNSTAWPLSLEPD
						PGMASACTTMHTTTIAEPDPG
						MSGWPDGRMETSSPTIMDIVVI
						*CAIAAEAIVLVSLLFVMLRYM
		İ				YRHMGTYHTNEAKGTGVADS
2 2 2 2 2		 	2.1-2	100		ADACPAGDPA
25333	55701	Α	25473	139	546	RPGMWSTRSPNSTAWPLSLEPD
						PGMASASTTMHNYHHCRSLIPG
1						M\SGWPDGRMETSTPTIMDIVVI
						AGVIAAV\AIV\LVSLLFVMVAL
						TCNRA/HRGTYHTNEAKG\TEF
						AESADAALQG\DPAL\QDAGDS SRK\EYFI
25334	55702	C	25474	142	408	SMX/E I F1
43334	20/02	л~_	1234/4	144	1700	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
2	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25335	55703	A	25475	44	727	ARAGSRSRGLPFCGTAGPPPKA
						TLWGGLLRLGSLLSLSCLALSV
						LLLAQLSDAANNFEDVRCKCIC
		l				PPYKENSGHIYNKNISQKDCDC
ļ	}			İ		LHVV\DPMPVPGA\NVEAYCLR
						CECKYEE\RSS\STIKVTIIIYLSIL
				İ		GLLL\LYMVYLTLVEPILKRRLF
		ŀ				GHAQLIQSDDDIGDHQPFGKCT
	1					RCASPLPQFEPTCWNKVEYAQ
						QALESFQVQEQRKSV\FDRACC
				İ		PQLNLGN
25336	55704	A	25476	11	1323	VVLLVSVNTEAEESEKAATEQR
		ĺ				/PLEGTDQTLDVEEEQEESKAA
						ACGSKKRVVPGIVYLGHIPPHF
						QPLHVRNLLSAYG/RSPSHYDL
						WNLKYLHHFTWSHHSEHLAFE
						RQVC\HRQHLRVEVTQAKCETI
						FYLQSVEWGQCFLAVDGDLAH
						PDGSWTFAQRPTEQELRARKA
						AQPVGKVLNYYPTQLSLECEVP
						SQPMGTGHSHRGLCQDVKVIN
						VPVSFVWCALVAGLLVVVPPPP
						PQAPEAEVSDISLKRGLQREIEQ
						CRRDIQNFPFPCASEELASQLFP
						LMEVPQGRGTTGFMNAPLTTS
						EVRGLKKELKPLLDDPERVAEQ
		1				IDQFLGSKLYTWTELMSILGILF
						SKEERNMICRAAMGAWERDYP
						AGQNIPAADVTFPARDPQWNN
	ļ					NNAAHQKNMRDLQELIIKGIKE
	}			1		SAPRPQNLTKAFDEQQEKDEGP
25337	55705	Α	25477	1	490	GTSGTRPLVSVNMEAEESEKA\
	}					ATEQEPLEGTEQTLYAEEEQEE
						SEEAACGSKKRVVPGIVYLGHI
						PPRFRPLHVRNLLSAYGEVGRV
						FFQAEDRFVRRKKKAAAAAGG
						KKRSYTKDYTEGWVEV\RDTR\
						IAKRVA\AALLK*PMGARRRSP
						FRYDLWNLKYLHR
25338	55706	A	25478	2	574	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	l • · · ·
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25339	55707	A	25479	160	1548	KTSSQTLSLALTMFQRLNKMFV
23337	33707		23477	100	1540	GEVSSSSNQEPEFNEKEDDEWI
		1				LVDFIDTCTGFSAEEEEEEEDIS
ŀ		l				EESPTEHPSVFSCLPASLECLAD
}		1				TSDSCFLQFESCPMEESWFITPP
ŀ		1				PCFTAGGLTTIKVETSPMENLLI
]				EHPSMSVYAVHNSCPGLSEATR
l						GTDELHSPSSPSILPNKMLSVHV
						GTAIHNQAALCQSVESYWQCGI
						QLLTQFQSQSKLGQLLAATCKE
						LPGPKESRRTAKDLWEVVVQIC
						SVSSQHKRGNDGRVSLIKQRES
						TLGIMYRYVLEPLVLTIILSLFV
						KLHNVREDIVNDITAEHISIWPS
						SIPKKQMSSRTFMSQEETVCRA
						AHTPSLPEEKWVEQLIYALVVA
						AARIMAMWGPNAAWLVFLAA
İ		1				AGPPPLTPPHSPP/PPPPPPALGR
						WMGPPLRDSASGLPRPPPGRISE
						DSGAWACGPALCPRPAIPSPAP
						ALRVPEALRLWGTCSV
25340	55708	A	25480	1	380	FTSPKTARNWGWLMQVRNQR
23340	33708	^	23400	['	360	LVLGKAGGKKGSPLEQRGGNG
						HHGGTRLICALV\LVS\MRPLAT
					ļ	FAEPQTETCTVAPRERQNCGFP/
						GVSRPPQCANKGCCFDDT\VRG
						VPWCFYPNTIDVPPEEECEF
25341	55709	A	25481	3	2367	VI WEI II NIID VII EELEEL
25341	55710	A	25482	2	1230	WRRSRRRRPPAPQAGGGDFRA
23342	33710	^	23402	2	1230	GISPPTGKKSLGLMEPQGNHYR
1						CRCLQVAGSQAWARCVSMPEL
						VVQKVVGHP/LVLLSVVDHFNR
						IGKVGIQKCVIGLFWGSWQKK
						VLDVLNSFAVPFDEDDKDNSM
						WFLDHDYLENMYGMFKKVNA
						RERIVGWYHTGLKIHKNGITIN
						ELMKRYYPNSVLIIIDVNPKDL
						GLPTEAYISVEEIQDDGTPTLKT
						FEHVTSEIGAESYLEKVATGKL
		1				RINHQIIYELQDVFNLLPDVSLQ
					1	EPIKAFYLRTNDQMVVVYLASL
						IHSVGLTDPFC\VALHNLINNKI
						ANQDGDKKEGQEKKES/KKDR
		1		1		KADKGKDKDKEKSDVKKEQK
						NHCLKEKYRFRQDSLPDDPRVL
				1		GLELTPPVARQYSPQDLSEIPRH
L	<u> </u>	L	i	<u> </u>	1	ADFRCDPAHFQLWWSQGETPS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25343	55711	A	25483	3	1042	DRYCCRCLRVAGSQAWRAGC
						VAMPELAVQKVVVHPLVLLSV
						VDHFNRIGKVGNQKRVVGVLL
						GSWQKKVLDVSNSFAVPFDED
		İ				DKDDSVWFLDHDYLENMYGM
						FKKVNARERIVGWYHTGPKLH
						KNDIAINELMKRYCPNSVLVIID
						VKPKDLGLPTEAYISVEEVHDD
8				1		GTPTSKTFEHVTSEIGAEEAEEV
						GVEHLLPDIKDTTVGTLSQRITN
		į]		HVHGLKGLNSKLLDIRSYLEKV
}						VTG\KLPINHQIIYQLQDVFNLL
						PDVSLQEFVKGFYLKTNDQMV
						VVYLASLIRSVVALHNLINNKI
						ANRDAEKKEGQEKEESKKDRK
						EDKEKDKDKEKSDVKKEEKKE
25344	55712	Α	25484	257	1566	GLEHLCNPRWPLLFLRAMDST
						VPSALELPQRLALNPRESPRSPE
						EEEPHLLS/SLAAVQTLANVIRP
						CYGPHGRQKFLVTMKGETVCT
						GCATAILRALELEHPAAWLLRE
						AAQTQAENSGDGTAFVVLLTE
		1				ALLEQAEQLLKFGLPRPQLR/EA
						YATATA/EEKQVGQL/AAAGIN
						VAVVLGEVDEETLTLADKYGI
						VVIQARSRMEIIYLSEVLDTPLL
		İ				PRLLPPQRPGKCQRVYRQELGD
						GLAVVFEWECTGTPALTVVLR
						GATTQGLRSAEQAVYHSIDAYF
				′		QPCQDPRLIPGAGATEMALAK
						MLSDKGSRLEGPNGPAFLAFAR
		ŀ				ALKYLPKTLAENAGLAVSDVV
						AEMSGVHQGGNLLMGVGAEGI
						INVAQEGVWDTLIVKAQGFRA
						VAEVVLQLVTVDEIVVAKKSPT
						HQQIWNPDSKKTKKRPPPVEK

SEQ ID	1 -	l .	SEQ ID NO:	l .		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	of peptide sequence	detetion, (=possible nucleotide insertion)
	<u></u>					
25345	55713	Α	25485	145	1918	PLEYFQRSVSGVPQLLFLEHLC
						NPRWPLLSLRAMDSTVPSALEL
						PQRLALNPRESPRSPEEEEPHLL
1					1	SSLAAVQTLASVIRPCYGPHGR
1						QKFLVTMKGETVCTGCATAILR
						ALELEHPAAWLLREAGQTQAE
			ŀ			NSGDGTAFVVLLTEALLEQAEQ
			:			LLKAGLPRPQLREAYATATAEV
						LATLPSLAIQSLGPLEDPSWALH
ŀ						SVMNTHTLSPMDHLTKLVAHA
					1	CWAIKELDGSFKPERVGVCALP
		İ	:			GGTLEDSCLLPGLAISGKLCGQ
			:			MATVLSGARVALFACPFGPAH
			:			PNAPATARLSSPADLAQFSKGS
						DQLLEKQVGQLAAAGINVAVV
		ŀ				LGEVDEETLTLADKYGIVVIQA
			:			RSWMEIIYLSEVLDTPLLPRLLP
						PQRPGKCQRVYRQELGDGLAV
				<u> </u>		VFEWECTGTPALTVVLRGATT
						QGLRSAEQAVYHGIDAYFQLC
						QDPRLIPGAGATEMALAKMLS
						DKGSRLEGPSGPAFLAFAWAL
				-		KYLPKTLAENAGLAVSDVMAE
						MSGVHQGGNLLMGVGTEGIIN
				•		VAQEGVWDTLIVKAQGFRAVA
				:		EVVLQLVTVDEIVVAKKSPTHQ
				•		EIWNPDSKKAK\KHPPPVETKKI
25346	55714	В	25486	1	642	DIWIN DOMARKATI II V DIKKI
25347	55715	A	25487	3	395	FLPVNLTHDWQGSSALATLERP
23317	33713	l' `	25 107	ľ		LSQVRLKRFVGTLIAFMVSAIVI
						LATASMAVASITESVQTAAFID
						NMARNVSNKLLLQQGIDQKIL
						ALLQSLEAALEYVGEQQDALA
						FQQQ*NCNWGHKHICIISLPWN
25348	55716	A	25488	3	358	HEPMCGETHQALQGAMEKLW
23310	33710	` `	25100	ľ		SRFMELMQE/KKVDLKERVEEL
						EHCCI*LSGERQTPSGSGKPGHG
		İ	1			KESCTLVRGAPASESCPPAGEYI
				1		ALYQSQRAVRKEEECISRLAQD
}]		KGEVKVKLLE
25349	55717	Α	25489	1	250	VQCGGIPPGQNNKEEMEVLPEP
22347	33111	l'^	23407	l'	250	PPPINRKKDKSYATAMGPFLRQ
						EALSGELLACLVIQD*QGNWV
						YKPISFNIYKKLRKSIRG
25350	55718	A	25490	1	702	I INTISTINI I KALANSIKU
25350	55719	A	25490	1	252	
25351	55720	A	25491	1	122	LAPLVTYFSSLSLGFFNLKGKL
عدوده	33720	^	23792	1	144	APSSWRC*GLNEIIRGKH
25353	55721	A	25493	3	505	711 SUNICO GENERIKOKII
23333	133,21	1.7	25795	l-'	505	

SEQ ID			SEQ ID NO:	L	1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		02/340,217	sequence	or peptiae sequence	deterion, i possible adelectide insertion,
25354	55722	Α	25494	1	3345	MRVGFRDIVGSLSEYSFGCPVN
						YEVFHSVWITILCCLMANDRKP
						QFLYFFQFFSFHGGRNLMSSFH
		1				PGNGHNDSRRSVFYANEKWEF
		1				LDPTPKDLEESIVQEEKKEPTPE
		1				GNKVSQRLKNLPHTSLKAKQL
						LQTSSTNRGTVITATRLAFDLRT
		1				QSTKCSHPASGSSGHSTDFQLE
	1					GEWGLWDLGTKGFKLPYSLSI
		1				DIDNVKSPHFPHELKTLTVSLGS
						NRRMGLEKKECHQKQGALREQ
		<u> </u>				LQVHIQNVEILVSE
25355	55723	Α	25495	296	1350	TLWLQGEQPSTQEIHEKV/LNE
						AMGAMMYHTIILTREDLEKFK
						APRVIVQIGSGYDNVDIKAASE
						LGIAVCNIPSAAVEETPDSTICHI
						LNLYQRNTWLYQALRGQAVA
						VRAKAFGFSVIFYDPYLQDGIE
		l				RSLGVQRDLLYQSDCVSLHCN
						LNEHNHHLINDFTIKQMRQGAF
]	LVNAPHGGLVDKKALAQALKE
]						GRIRGAALYVRESEPFSFAQGP
					[LKDAPNLICTPHTAWYSKQASL
					}	EMREAAATEIRRAITGHIPESLR
						NCVNKEFSVTSAPWSVIDQQAI
						HPELNGATYRYPPGIVGVAPGG
		ŀ				LPAAVEGIIPGGIPVTHNLPTVA HPSQAPSPNQPTKHGDNREHSN
25356	55724	A	25496	694	1185	PKPFKEGTIRGAALDMHESEPF
23330	33724	^	23490	1054	1105	SFAQGPLKNAPNLICTPHTAWY
İ		l		ļ		SEQASLEMREAAATEIRRAITG
						CIPESLRNCVNKEFFVTSAPWS
		l				VIDQQAIHPELNGATYRYPPGV
						VGVAPGGLSAAMEGIIPGGIPV
						THNLPTVAHPSQAPSPNQPTK*
						MK*KQMGEVA*GAARIRGETL
						GLNGFGCTGQA/VAVRAKAFG
						FSVIFYDPYLQDVVERSLGVQR
						VYTLQDLLYQSDCVPLHCNVN
						EHNNHLINDFTIKQMRQGAFLP
				1		PKPFKEGTIRGAALDMHESEPF
						SFAQGPLKNAPNLICTPHTAWY
]		SEQASLEMREAAATEIRRAITG
						CIPESLRNCVNKEFFVTSAPWS
						VIDQQAIHPELNGATYRYPPGV
						VGVAPGGLSAAMEGIIPGGIPV
						THNLPTVAHPSQAPSPNQPTKH
L	<u> </u>		L	L	1	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25357	55725	I _A	25497	290	1251	 PFMLYVSISTLYIAFCHILNLYR
23337	33723	ļ'`	25177	2,0	1231	RNTWLYQALWEGTRVQSVEQ
						MGEVA*GAARIRGETLGLNGF
						GCTGQA/VAVRAKAFGFSVIFY
						DPYLQDVVERSLGVQRVYTLQ
						DLLYQSDCVPLHCNVNEHNNH
	1					LINDFTIKQMRQGAFLVNAARG
						GLVDEKALAQALKEGTIRGAA
	}					LDMHESEPFSFAQGPLKNAPNL
1						ICTPHTAWYSEQASLEMREAA
	1					ATEIRRAITGCIPESLRNCVNKE
		1				FFVTSAPWSVIDQQAIHPELNG ATYRYPPGVVGVAPGGLSAAM
						EGIIPGGIPVTHNLPTVAHPSQA
25358	55726	-	25498	1	1520	PSPNQPTKHVDNREHPNEQ
23338	55726	A	23498		1538	MLLQKTQFEYAFKKKRKLEES
						KKRGAGPASSDPRWRGSSQTA
						PHASEVSGAPLGRSCETTRDTG
						VGPSPRQWLGAGAGTWQHRA
						RAASTPSSRGLRALQTWGGGE
						VMNGPLH\PPPPRVALLD/SRNC
				1		TVEMPIPKDLATMAFCDTHSTQ
						EIHEKILHEALGAMRYHTITLA
		1				RADLEKFKALGVIVRIGSGYDN
			ľ			VNIKAACELEIAVCNIPSAAVEE
						TANSTICHILSLYRRNTWLCQA
						WARKHVCYSMGQIREVALGG
·						AHIGGETQG\LIGFGSTGQAVPV
						RAEAFGFSVIFYDPYLQDGIERS
	İ	1	·			LGVQRMSQGAFLVNAARGALV
						DEKALAQALNNGKIR/GAALDV
	1					PQSELFSFAQG\SLEMREAAATE
						IRRTITGCIPESLRNCVNKEFFVT
						SAPWSVIDQQAIHPELNGATYR
	İ					YLPGFVCVAPGGLPAATEGIIPG
						GIPVTHNLPTVAHPSQAPSPNQP
		1				TKHGDNPEHPNEQSRECRKVII
						QIHLRPRDSEKLMNSEKTNLTV
	1	1.	0.5555		2505	FLADSGHMHY
25359	55727	A	25500	1	2595	CCHD A V DEGIGE V D VICE LA SUB-
25360	55728	Α	25501	561	1023	SGHDAYREQIQEYRVISLLLNP
						N\DHVNKCQSTNDAYPTGFRIA
						VYSSLIKLVDAINQLREGFERK
]			AVEFQDILKMGRTQLQDAVPM
1						TLGQEFRAFSILLKEEVKNIQRT
1		1			:	AELLLEVNLGATAIGTGLNTPK
						EYSPLAVKKLALKSLAFVGNLF
25361	55729	A	25502	2	227	
25362	55730	Α	25503	262	428	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25363	55731	A	25504	389	2310	ATSGPPGRSLGSACTIWVLVWT
						GSIPIYSFHGKVEKYMCFHNMS
	·					DDTWSAKVFFPLEVFGFLLPMG
						IMGFCCSRSIHILLGRRDHTQD
						WVQQKACIYSIAASLAVFVVSF
						LPVHLGFFLQFLVRNSFIVECRA
	Ì					KQSISFFLQLSMCFSNVNCCLD
						VFCYYFVIKEFRMNIRAHRPSR
						ATEEVSKNLVAMKEILYGTNE
						KEPQTEAVAQLAQELYNSGLLS
		1				TLVADLQLIDFEASLGLHGLEQ
	•					AVDLALPIAQAQGEVLNNGKC
						MDQFPVDVYQGGAGT\YVNMS
			•			TNEVLANIGLGLMGHQKGEYQ
		l				YLNPNDHVNKCQSTNDAYPTG
						FRIAVYSSLIKLVDAINQLREGF
						ERKAVEFQDILKMGRTQLQDA
						VPMTLG\QEFRAFSILLKEEVKN
						IQRTAELLLEVNLGATAIGTGL
						NTPKEYSPLAVKKLAEVTGFPC
						VPAEDLIEATSDCGAYVMVQG
						ALK\RLAVKMSQI\CNDLRLLSS
						GPPG\SSIMPAKVNPVVPEVVN
1						QVCFKVIGNDTTVTMAAEAGQ
		1				LQLNVMEPVIGQAMFESVHILT
ŀ	1	-				NACYNLLEKCINGITANKEVCE
						GYVYNSIGIVTYLNPFIGHHNG
į		1				DIVGKICAETGKSVREVVLERG
				1		LLTEAELDDIFSVQNLMHPAYK
25364	55732	Α	25505	133	375	
25365	55733	Α	25506	3	505	ARGREGVSRRERPLLSARRLGA
		1				EWDSAGAETASGLQVPRLGGS
						WPQTSVFSAAVKARGCRIATLP
						MDLIGFGYAALVTFGSIFGYKR
						RGGVPSLIAGLFVGCLAGYGAY
ļ						RV\SNDKRECKKCHWFTAFLPG
		1				LPSMGVRFKRSKKIM\PAGLVA
		L				GL\SLM\MILRLVLLLL
25366	55734	Α	25507	244	696	

SEQ ID NO:	SEQ ID NO: of peptide sequence	•	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25367	55735	A	25508		1442	MGQLHQAQASADIATSRLQQT SLTDVTRHQQFFSTYNTDDFIQ FTATYREQTRTRRQRQYAVNH VTFFFGKCTFRDRHRHRVIGIVL NIGLTFPAHQTQDGIGSTRSQR QSRLVANKVATRNLVKHFDDD REANRGIQEAQTEHHHRGVLV NESRQRFGCQQHNRHRNHYRR HHHRQMVNHPYCSNYRIQRED GVKDDNLCDDHPETGIQLYASL LLQVFFFAANIYGWYAWSRQT SQNEAELKIRWLPLPKALSWLA VCVVSIGLMTVFINPVFAFLTR VAVMIMQALGYLQGVMPELQA DAFPVWDSCMMVLSIVAMILM TRKYVENWLLWVIINVISVVIF ALQGVYAMSLEYIILTFIALNGS RMWINSALIHQVVLDLIQQGIVI VWRDHLHMQRLHRLLAYHPC MYLVHIADFRDFPAHTRLQVQ RFPAIPVHLQVIHSRYLSTVATR YAESEATPVSGALGLLCPTLPIV
25368	55736	A	25509	133	375	TAESLATI VSGALGLECT TETTV
25369	55737	A	25510	2	471	AFAQRETAGRRVGQRWCGDCF RTPGTALGGSWPQTSVFSAAV KARGCRIATLPMDLIGFGYAAL VTFGSIFGYKRRGGVPSLIAGLF VGCLAGYGAYRV\SNDKRECK KCHWFTAFLPGLPSMGVRFKRS KKIM\PAGLVAGL\SLM\MILRL VLLLL
25370	55738	A	25511	3	432	NSRVDDFVAAQDAKGKKVAP APAVVKKQEAKKVVNPLFEKR PKNFGIGQ\QRLLARAEKKAAG KGDVPTKRPPVLRAGVNTVTT LVENKKAQLVVIAHDVDPIELV VFLPALCRKMGVPYCIIKGKAR LGRLVHRKTCTTVAFT
25371	55739	A	25512	59	882	SRLKNASIFVASPSGKKAKGKK VAP/APAVVKKQEVKKVVNPLF EKRPKNFDIGQDIQPKRDLTSFV KWPPLYQRQRAILCKWLKVPP EINQFTQALDHQTAALLLQLAH KYRSETKQEKKQRLLALAKKK AAGKGGIPTKRPPVLRAGVNTI TTLLENKKAQPVVIAHDVDPIE LVVFLPALCHKMRVPYCIIKGE ARLGRLVHRKTCTTVAFIQVNS EDKGALAKLVGAIRTDYNDRY NEIRRHWGGNVLGPKSVACIG KLEKAKAQELATKLG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25372	55740	A	25513		833	GTRPKMPKGKKAKGKKVAPAP AVVKKQEAKKVVNPLFEKRPK NFGIGQDIQPKRDLTRFVKWPR YIRLQRQRAILYKRLKVPPAINQ FTQALDRQTATQLLKLAHKYR PETKHEKKQRLLARAEKKAAG KG\DVP\TKRPPV\LRAGVNTFT PL\VENKKAQLVVIAH\DVDPM RGWLSFLPAL\CRKMGVPYCIIK GK\ARLGRLV\HRKTCTTV\AFT QVNSESQRRLWAKLVEAIRTN YND\RYDEIRRHWGGNDLRPK\
25373	55741	A	25514		1176	SVARI\AKLEKAKAK\ELATKLG MEKELVDHTGLFLSAHASEAL ALKLMIYSEVAHKLRAVRSQH PDMGVSLPLTPANSCPKTIERM RRSRELEREKDQSIETDHLPWA FGSTFLKTPLSCKTLMKYICYA FIINLSFVKGVSAVTLMMSEES NMQMQFSVSITSGFGPFKSHIIA AIPNAERKESQGEKMAPAPAVI KKQEAKEVVNPLFEKRLKNFGI GQDIQPQRDLTRFVKWPHYIRL QRQRAILS\AHKYRPETKQEKK QRLLAQAKKKATSK/EDIPTKR PPVLRAGVNMVTTLVENKKAQ LVMTAHDVDPIELAVFCLPAPC CKKGVPYCIIKGKARLGHLVHR KTCTTVVFTKVNSEDKGALAK LVEAIRTNYDRYDEIHCHWGG NVLGLKSVAHIIKLEKAKVKEL
25374	55742	A	25515	2	524	
25375	55743	A	25516	144	412	AGSCPFAAGPGPLGSGRCFVLIR FILIQNRAGKTRLAKW\YMQFD DDEKQKLI\EEVHAVVTVRDAK HTNFVGVSGTFKIILPPLWLASN
25376	55744	Α	25517	38	476	
25377	55745	A	25518	235	892	RSRVGTNRLFGETYPRASPGPQ STALLTAYKKMTDLVAVWDV ALSDGVHKIEFEHGTTSGK/REV VYVDGKEEIRKEWMFKLVGKE TFYVGAAKTKS/ATINIDAISGF AYEYT\LEINGKSLKK\YMEDRF KNPPILWVLHM\DG\ENFRIVLE KDAMDVWCNGKKLETARDFC KMSDTISDDTKVDEQERALSRT PEDKWNFRPAVPMRGERGSPS HCGQAH

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence	ł	,	sequence	,	
		<u> </u>				
25378	55746	Α	25519	1	1233	MQLKFPGRQKIHISKKWGFTKF
		Ì				NADEFEDMVAEKRLIPDGCGV
					į	KYIPNRGPLDKCHHGKCVATG
						TSYMCKCAEGYGGDLCDNKN
						DSANACSAFKCHHGQCHISDQ
		1				GEPYCLCQPGFSGEHCQQGLLG
						GVREMLRCKVRSNIVTMKAKL
			ŀ			SFRICMTLTILANMWPRPILGG
1						VDQIPIKPKSKVLYLGAASGTT
						VSHVSDIIGPDGLVYAVEFSHR
						AGRDLVNVAKKRTNIIPVLEDA
		1				RHP/HQPDQ\ARSVALNAHTFLR
		1				NGGHFLISIKANCIDSTASAEAV
		1				FASEVRKLQQENLKPQEQLTLE
		1				PYERDHAVVDYSSSATEQRRT
į						ENDSDKLTEVGFRKLVINFSKL
						KEDVRTHHKEVKNLEKRLDEW
	ľ					LTRINSVEKTLNDLMEAENHGT
1						KKYVTQAQASVAYLIKWKKG
25379	55747	A	25520	78	978	AAGHSAKKSCLSLLPAWSFPLG
						STRAQWHVRRLQARISGGGRH
						RCGRRGQ\GKTRRRMGPMVVS
						VEPHRHEGVFIYRGAEDALVTL
	İ					NMVPGQSVYGERRVTVTEGGV
1						KQEYRTWNPFRSKLAAAILGG
						VDQIHIKPKSKVLYLGAASGTT
						VSHVSDIIGPDGLVYAVEFSHR
						AGRDLVNVAKKRTNIIPVLEDA
						RHPLKYRMLIGMVDVIFADVA
						QPDQSRIVALNAHTFLRNGGHF
		1				LISIKANCIDSTASA\EAVFASEG
						EKLQQENLKPQEQLTLEPYERD
						HACV\VGVYRPLPKSSSK
25380	55748	$\frac{1}{A}$	25521	230	314	QMPGNNPAFSA*CDAGASYQA
23380	33748	A	23321	230	314	YVNSAIY
25291	55740	A	25522	1360	1848	RFRALV/TEGFSLRELFPVATVL
25381	55749	A	23322	1300	1040	VASST\VTKRFIFLLAAD\VRLA
						LRRSI\THPFVRKQELTVYTLNN
					,	ELENLLTNVVNQAQQGGKVML
						DSVPVDPNMLNQFQSTMPQVK
				1		1
						EQMKAAGKDPVLLVPPQLRPL
				1		LARYARLFAPGLHVLSYNEVPD
		<u>L</u>		<u></u>		ELELKIMGALM

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25382	55750	A	25523	91	1184	AAACSPPPPARLTYGSETLRGG DCSKMAAGCCGVKKQKLSSSP PSGSGGGGGASSSSHCSGESQC RAGELGLGGAGTRLNGLGGLT GGGSGSGCTLSPPQGCGGGGG GIALSPPPSCGVGTL\LSTPAAA TSSSPSSSS/APASSSSPGSRK\M VVSTE\MC\CFCFDVLYCHLYGI \QQP\RPPRFTNEPYPLFVTWKIG RDKRLRGCIGTFSAMNLHSGLR EYTLTSALKDSRFPPMTRDELP RLFCSVSLLTNFEDVCDYLDWE VGVH\GIRIEIHQCKKASKR\TA TYLPEV\AKEQGWDHIQTIDSLL RKGGYKAPITNEFRKTIKLTRY RSEKMTLSYAEYLAHRQHHHF QNGIGDPLPAYNHYS
25383	55751	Α	25524	3	154	Q.verez. 2
25384	55752	A	25525	575	1707	TALLLTQSLFGGLFTQTHMKFG AVTRI*NRKTGNSKKQSTSPPPK ERSSSPATEKSWMENDFDELRE EGFRRSNYSELREDIQTKGKEV ENFERNLEECITRITNTEKCLKE LMELKTKARELCEECRSLRSRC DQLEDCLLKKLHSHQANPEFDL ITEHLTLCIRISIQIKAYLSLSKA QSSLSHRPPTYLAYLFNGFRLF KNLKAYLHPGEINSFIAHTKPV WWSLHTDAHEIWCRDSDLKLV PTLPLIPLEAALRNITHSLSIPPP KNFRRPNTSTLFCVIFLINIRRQE YQASEPKPSHRIPCDLQRIKYLG IQLTRDVKDLFKENYKPLRNEI KEDTNKWKNIPCSWVGRINIVK MAILPKPLLVIPRQTGSGMDPQ QTSGDLQKCLTVLRKTNKQKAI ASTSRKRMTMQKLHLKEKLINS KDQRTNDKNHMIISIDAEKAFN KTEHRFMLKTLNKLEINSRWIK DLNVKPKTIKTLEENLGNTIQDI VMDKDFMTKTPKAMATKAKI DKWDLIKLKSFCTAKETIIRVY RQPTERESIFVIYPSDKGLISRIY KELAIPGKGNSSNVSKDSKMSK HQKMQGENKHD

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25385	55753	A	25526	110	569	TALLLTQSLFGGLFTRTRMKFG AMTRIG\DLPWEINPLSSCSLLC EKHPPTTSGPQTDQPKKHLTNF KSETKETCFIHEPKTLAPVTDW EGSLPLVFNHCRDGLSDHSATF QGCQTMQGCLPWSFTLSGKSR FSGEGANHHGHRASGNSHSGS
25386	55754	A	25527	2	,	LFGGLFTRRRMKFGAVTRIR\DF PWEINPLSSCSLL/CEKDPRTTS GPQTDQPSKHLTNLKSASTPPP YNPFITSPPHTRSGLQFRSTSSPP APAQQFTLKKVAEAKGIVKVN APFSLSDLSQISVRLGSFIKYEKS SPVHGSFGSNPETLYSPRP
25387	55755	Α	25528	162	206	
25388	55756	A	25529	74	446	TAMLLTQSLFGGLFTWTRMKF GAVTRIG\DLPWEINPLYSCSLL SRFKKIKACYHSPATSWPFKTY KLSLQLPHFTCPKTRQALQVSS GAVPYQPNCFAYPPCGAQPVPS FVLNTFLHNSLCRA
25389	55757	Α	25530	66	311	FGAVTRIG\DLPWEINPLSSCSLL REKEPPTTSGPQTNQAKKHLTN FKSGACYTCRKSGHWAKECPQ PGIPPKPRPICVGPH
25390	55758	A	25531	728	825	MKFGAVTRIG\DLPWEINPRSSC SLLHEKDPPM
25391	55759	A	25532	291	311	TALLLTQSLFGGLFTQTHMKFG AVTRI*NKKNNTK
25392	55760	Α	25533	1	1056	
25393	55761	A	25534	245	487	TALLLTQSLFGGLFTQTRMKFG AVTRIG\DHPWEINPLSSCSLFQ EKDPPTT/SGPQTDQPKKHLTNF KSTSFVFSSCIPPP

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	i e	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25394	55762	Α	25535	818	2391	TALLLTQSLFGGLFTQTRMKFG AVTRIG\DIPWEINPPSSCSLFLP/ RLPLLHFPGAGPSTPQPLL/HSPS AASPAFLGQGQDLINLVFKVYN NRKKLQFLASTVTQTPATSPAH KNFQTPEPQQPGVPPEPPSPGAC YKCQKSGHQAKECPQLRIPHLS YAWTFNDNPLYVQEDNRRFVS QETGNLYIAKVEPSDVGNYTCF ITNKEAQRSVQGPPTPLVQRTD GVMGEYEPKIEVRFPETIQAAK DSSVKLECFALGKIDDSIREKG AIRIKPSSILMRCNCLVQSPILVG EGWTGARCQGKSSTANPKLSL KSRTSNKKMKAFMSALQATFE EETLQRVNSFFMHNVLDNCLLP NSGTCLFETAPPEWEQKIQNTH LSIYDNLLWECKASGKPNPWY TWLKNGERLNPEERIQIENGTLI ITMLNVSDSGVYQCAAENKYQI IYANAELRVLGSQLVHFRENGT CLSDEVIRDSDFMSGMTVVHAF TSGSQGTTTENQDELQIKFISAR
25395	55763	Α	25536	425	749	HSQATEGINSTSAKFD IASLLTQSLFGGLFTWTHMKFG AVTRT/RGDLPWEINPLSSCSLL HEKDPPTTSGPQTDQPKKHLTN FKSAQLKASGK/DLQKPYQHFQ TLQQLLPLQWAPCYGPSSVSI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25396	55764	A	25537	2	2204	I IDMIFTPGPPSTPKHKKSQKGSA
23390	33704	Δ.	23337	2	2204	FTFPSQQSPRNEPYVARPSTSEI
	}					EDQSMMGKFVKVERQVQDMG
						KKLDFLVDMHMQHMERLQVQ
						VTEYYPTKGTSSPAEAEKKEDN
						RYSDLKTIICNYSETGPPEPPYSF
	1					HQVTIDKVSPYGFFAHDPVNLP
						RGGPSSGKVQATPPSSATTYVE
						RPTVLPILTLLDSRVSCHSQADL
						QGPYSDRISPRQRRSITRDSDTP
,						LSLMSVNHEELERSPSGFSISQD
	1					RDDYVFGPNGGSSWMREKRYL
						AEGETDTDTDPFTPSGSMPLSST
						GDGISDSTVPLFLSSEILQKQVG
						QSITSMLGFLSRGPSMKLCMGL
						ACVLSLWNTVSGIKGEAKKEK
						GMTFLPTTDSKKFFSLLSVTSYS
						SFAFHKFSVAVYNISNLKTVDP
						AKFPTRYCYCLNNRTNDLSDFT
						ALLVDIIGNSTSYLTEIFKSTSIL
						SVNQSNESDCIFICVMTGKSGR
						NLSDFWEIEEKYPIINYTFTSGL
						SGVLALLLTQSLFGGLFTRTRM
					İ	KFGAVTRIG\DLPWEINP/LSSCS
						LLHEKDPPTTSGPQTDQPKKHL
						TNFKSAARPTFLGQGQVPLNPF
						SFTLSGKSCFPRRQEPPNRLFPH
						PNLLSLCPNPLFLCPNPFSTFLE
				1		GKNHHPHPFSVSQVFSLGLPPS
						LWEHSPMAHRLRVQPRQQPPDI
						HKQVISLQSGHSLLVKSQSWSQ
						DPIKFQLPHHHKNTASVCINPSV
25397	55765	A	25538	110	542	TALLLTQSLFGGLFTRTRMKFG
2337	ده ۱ دور	^ <u></u>	0502	110	J-12	AMTRIG\DLPWEINPLSSCSLLC
				[EKHPPTTSGPQTDQPKKHLTNF
			!			KSETKETCFIHEPKTLAPVTDW
						EGSLPLVFNHCRDGLSDHSATF
						QGCQTMQGCLPWSFTLSGKSR
						• •
25200	55766	 	25520	1	781	FSGEGASCSLPG MPLQPLATPFTSPGSPSVAQVT
25398	55766	Α	25539	1	/01	
				1		TNTGEPGFAQQEYAETPQTTAR
						PPHPHPALRALQASVHQTPRLG
				1		DCLGSSESATALHGSGEPNKLA
						PPGRPTERGGLFKCSVPLRTVR
						GCRAGLRGTETVPSSLQRQPAG
						DVGWINNKSTTALDRTPTDTQ
						RTRRPRPPKQKQRGRALKERPR
						DSDSAREGPLKAVAGLFGGLFT
						RTCMKFGAVTRIG\DLPWEINPL
						SSCSLLHEKDPPTTSGPQTDQPK
L	<u> </u>	L		<u> </u>		KHLTNFKSDLLSLATED

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
25399	55767	Α	25540	1571	1781	TALLLTQSLFGGLFTRTRMKFG
				·		AVTRIG\DLPWEINPSSSCSLLRE
]				KDPPTTSGPQTDQPKKHLTNFK
						SA
25400	55768	Α	25541	2150	3195	TVLLLTQSLFGGLFIWTRMKFG
		1				AVTRIG\DLPWEINPLSSCSLLH
		1	i i			EKDPPTTSGPQTDQPKEHLTNF
						KSDED
25401	55769	Α	25542	4688	5149	TAMLLTQSLFGGLFTRTRMKFG
		l				AVTR/IRGTDLPWE\INPLSSCSL
						LREKDPPMTSGPQTDQPKKH\L
						TNFKSGACYTCWKSGHWAKER
						PQPGIPPKPRPICVGPHLKIRTCS
		1				THLGQATPRAPGTLAQGSLTDS
		1				FPDLLGLAAEDSHCLIASEAP
25402	55770	Α	25543	1	433	MAAFPAALNRPTRARARRRIS
		1				AHSAASAAAAVVVAGTARGM
						VPGSEGP\CRA\GDEVADWEIVI
						EGTANLGPYFEGLRKHYLLPAI
						EYFNGGPPAETDFGGDYGGTQ
						YSLVVFNTVDCAPESYVQCHA
		<u> </u>				PTSSAYEFVTWLDGIK
25403	55771	Α	25544	1	235	
25404	55772	A	25545	1	301	
25405	55773	Α	25546	2	438	
25406	55774	Α	25547	1058	1494	KSVRLVVN/YLRTQKAVVRVSP
						EVPLQNILPVICAKCEVSPEHVV
						LLRDNIAGEELELSKSLNELGIK
					ļ	ELYAWDNRRVPSAVPMQGPWP
						AEHFLQVNEQPPAGSPSPCPSSS
						SSSSGLRGGMQTANGPACALTP
						TCAQTDCPCPTSH
25407	55775	Α	25548	[1	3965	RRFQPLPSAALRRCPPPARVGG
						RWVSAEKVNEGGSGGGGSQRL
		1				TRRARSTARVPAPGTMDAPRAS
		l				AAKPPTGRKMKARAPPPPGKA
		1				ATLHVHSDQKPPHDGALGSQQ
		1				NLVRMKEALRASTMDVTVVLP
		1				SGLEKRSVLNGSHAMMDLLVE
						LCLQNHLNPSHHALEIRSSETQ
					1	QPLSFKPNTLIGTLNVHTVFLKE
					:	KVPEEKVKPGPPKVPEKSVRLV
1			İ			VNYLRTQKAVVRVSPEVPLQNI
	<u></u>	<u>L</u> .				LPVICAKCEVSPEHVVL
25408	55776	Α	25549	1	388	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25409	55777	A	25550	1	1748	MVLARVARRPAGSAEAQASLA
23409	33111	A	23330	1	11740	GKAALPERARKMSSPGIDGDPK
						PPCLPRNGLVKLPGQPNGLGAA
						SITKGTPATKNRPCQPPPPPTLP
						PPSLAAPLSRAALAGGPCTPAG
İ						GPASALAPGHPAERPPLATDEK
						ILNGLFWYFSACEK\CVLAQVC
			•			KAWRRVLYQPKFWAGLTPVLH
						AKELYNVLPGGEKEFVNLQGF
1						TARGFEGFCL\VGVSDLDICEFI
1						DNYALSKKGVKAMSLKRSTIT
						DAGLEVMLE\QMQGVVRLELS
						GCNDF\TEAGLWSSLSARITSLS
						VSDCINVADDAIAAISQLLPNLA
1						ELSLQAYHVTDTALAYFTARQ
				İ		GHSTHTLRLLSCWEITNHGVVN
						VVHSLPNLTALSLSGCSKVTDD
		1				GVELVAENLRKLRSLDLSWCP
		l				RITDMALEYVACDLHRLEELVL
		1				DRCVRITDTGLSYLSTMSSLRSL
i						YLRWCCQVQDFGLKHLLALGS
						LRLLSLAGETPVSALTLAVTTH
		l				PHLVHPPNLPGPCANSHPARTH
	:					SPGSEAEEEGGAARPSPPEPWA
		l				ARTEPPSDPAAGCPLLTTTGLS
		l				GLVQLQELEELELTNCPGATPE
	{	ŀ				LFKYFSQHLPRCLVIE
25410	55778	A	25551	1	377	MRRLTRRLVLPVFGVLWI/DGA
25410	33770	ľ` `	25551	[*		AVLLGNQEEVGGAD/WDLKCR
				}		PL/EVNVSKSDDTLKINGVEDH
				-		KTIFDGDGKTYQNVQQFIDEGN
						YTSGDNHTL/SRPSLR*RQRP*IP
		1				CIRSQHGNRKRIPRRRIFI
25411	55779	A	25552	222	287	Chesqueria indui i
25412	55780	A	25553	2	918	GHCEVNRDWLQPLLHRVKEDY
23412	33780		25555	2	710	TRVVCPVIDIINLDTFTYIESASE
	1					LRGGFDWSLHFQWEQLSPEQK
İ						ARRLDPTEPIRTPIIAGGLFVIDK
						AWFDYLGKYDMDMDIWGGEN
						FEISFRVWMCGGSLEIVPCSRV
						1
						GHVFRKKHPYVFPDGNANT\YI
						KNTKRTAEVWMDEYKQYYYA
						ARPFALERPFGNVESRLDLRKN
						LRCQSFKWYLENIYPELSIPKES
						SIQKGNIRQRQKCLGISKGRTTK
						KPQT*S*A\LCQGQRRRCQSPRV
						WGLSHYTQADSFQGELVALFSS
		<u> </u>		1		FTFVSLGAPVVSCSFARMGD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25413	55781	Α	25554	84	2804	PRGSWRCRRDLKCRPLSHSDA
						DWDDLWDQFDERRYLNAKKW
						RVGDDPYKPVCFSTSGRVSGSP
		Ì				AIRAIPDTRHLRKKATSNSLSRQ
		ŀ				QLASPLPRSSYCGVTPGPIAIGP
						LWTSGSLRTPNHIDTVFAPGLP
Į						QPWFHLNYLRAMPSLSSAPKCT
		ļ		}		LLVYCTDLPPTSIIITFHNEARST
						LLRTIRRPPAVSTSQMAMTAAA
						LTPAPLRTYSVHLSCRAHQLVP
						MPQRKKYDETWVREHTKHSKI
						SLFPSCRLHFSCLSRRRMAVGS
						GKFQVMGDLGEPFLRVRWLRI
						WTLGPDFHGSALNSVLNR\TPT
		ļ				HLIREIILVDDFSNDPDDCKQLI
						KLPKVKCLRNNERQGLVRSRIR
		i				GADIAQGTTLTFLDSHCEVNRD
						WLQPLLHRVKEDYTRVVCPVI
1						DIINLDTFTYIESASELRGGFDW
						SLHFQWEQLSPEQKARRLDPTE
						PIRTPIIAGGLFVIDKAWFDYLG
						KYDMDMDIWGGENFEMMTVQ
						VLQPRRSEALICTWKVHHLTRE
						STALLIALVQAPFTGPQAKKLQ
						QSPCPAMLAPEYAPEASVLVCT
						VSLLTYTTLQSEISFRVWMCGG
		1				SLEIVPCSRVGHVFRKKHPYVF
						PDGNANTYIKNTKRTAEVWMD
						EYKQYYYAARPFALERPFGNV
						ESRLDLRKNLRCQSFKWYLENI
				-		YPELRYPSYLLELWLAESRGPQ
		1				LAWSGEETDNDQLDEGFYVYQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	1	
25414	55782	A	25555	1	2439	MPLKKKGDEPQMEKYIQERSM
25414	33762		23333	1	2437	LAPPSCSDHRDTHSRCIPHIQRL
						FQCPLGPWFQGEQRNQARGEG
						GALHPLPVTAGTVALPGAAQPT
						PARGNVTESLQGATVKLVCKK
						TSFIFIYCQHPAQCLALACDHEC
						IDLCTHVASTSLCLRGYQGMKR
						TIERALTATLAQGTAVPPFPSDV
						PSQKDAAVRGKGDGSEWERTL
						TMKRSNLNSEKILPAILLLTEFP
						RHQSPCSFDKPHHPNTLGCWLP
		İ	1		<u></u>	TVLETRGGTNGQSSGKREATLC
						PCPLLGLPISEEDCVPPALLLPPR
						G\LPPGLQLSWAPVPQFPPLGPS
		1				SLGIPA/LPLVLNLGCLPSLVCPP
						TSAHTSEVAPSLKSPAEPSPPAR
						LPRSHQPALSREMRHSDYSVSV
						HAICWDHDFYTLVTHYMNNTS
						PISSKRSSDKLNPHVLVVQDSM
						TILSLHNHLALAGQDMMVWEP
					1	RKTLQRRQQPDSSSHSPPPALV
		1				ECHPLEPEPLVAGIMYSFSGLQ
						GVGLRDRAVDETNKRNEKRTL
				1		EMSRVEALSSFENSEFGCPPSIN
						TREITGNSLEGNVKKGVSVPNC
						TAPGKRTEKKYSKIHEMVKHS
						GNCKRTDGIWEEYGWSWRLH
						MKKVFPGFMLVKGGEDAAPGT
						KEIKSVSGIFQRGKKKPRGRKE
						KNRKKRGFAFSKETNFVYLLIL
						VCLIGKVVVWLTSACAEKPKP
						HGEATCRHSGPSAPLNFOPTRV
25415	55783	A	25556	2	1178	AKCVEVCGKREPPSMSPLNQSA
23413	33783		25550		1170	EGLPQEASNRSLNATETSEAWD
		Į				PRTLOALKISLAVVLSVITLATV
		ł				LSNAFVLTTILLTRKLHTPANYL
						IGSLATTDLLVSILVMPISIAYTI
						THTWNFGQILCDIWLSSDITCCT
						ASILHLCVIALDRYWAITDALE
			İ			YSKRRTAGHAATMIAIVWAISI
]	CISIPPLFWRQAKAQEEMSDCL
						VNTSQISYTIYSTCGAFYIPSVLL
						IILYGRIYRAARNRILNPPSLYG
						KRFTTAHLITGSAGSSLCSLNSS
						LHEGHSHSAGSPLFFNH\VKIKL
						VTVALKRKEDSAARERKATKIL
		1				GIILGAFIICWLPFFVVSLVLPIC
						RDSWWIHPALFDFFTWVGYLN
						SLINPITYTVFNEEFRQAFQKIVP
						FRKAS
25416	55791	1	25557	3	146	I AKAS
25416	55784	A	23337	J ²	1140	<u> </u>

SEQ ID	SEQ ID NO:		1		1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25417	55785	A	25558	45	592	PSPPPPARAPSARTRRAPIPAAG QGRARPPRTVSRCGSAAMATKI
						DKEACRAAYNLVRDDGSAVIW
						VTIKYDGSTIVPGEQGAEYQHFI
						QQCTDDVRLFAFVRFTTGDAM
	İ					SKRSKFALITWIGENVSELQRA
						KTGTDETLVKEVVQNFANEYVI
						SDRKELEEEFIKS\ELKKAVGAN
						YDAQTE
25418	55786	Α	25559	249	1808	INGGTLTVTLSVTNQGAETLPF
						GTGWHPYFPLSPQTRIQAQASG
						YWLEREQWLAGEFCEQLPQEL
						DFNQPAPLPRQWVNNGFAGWN
						GQARIEQPQEGYAIIMETTPPAP
						CYFIFVSDPAFDKGYAFDFFCLE
						PMSHAPDDHHRPEGGDLIALAP
						GESTTSEMSLRAYNDTNRRSDK
		1				AFTPHPTCFPSNQNNSASPPTIT
						PDTATIAPVSHKAFAGNALRSII
						SDGKLTAGSVINNASAGAVPIP
						ASIIVCTIGISAAVGMTNKHPAT
						AIDTTHIMLLSTAPSAWGNNQT
						RAAPNTSLLAAYRFLIDSRDTE
						TDSRLDGLSDAFSVFRCHSIMN CVSVCPKGLNPTRAIGHIKSML
						LQLPPNVEKFAGAYLIQFALFQ
					ļ	PLTHNRLMFFGAFADFLHADQ
						TFPGQITAAVKGQFVRRRLRFF
						AIISPSRTLS/YHTP*PRR*PR*SN
						FVTGRDAGRGLYNRQ*KTS/PS
		1				SICSA*GANPVVHHAMTVKFA*
						WQPIMKISSTSFPSMRPLSHMK
						TINGLTTFVAW
25419	55787	В	25560	1	567	
25420	55788	A	25561	1	741	MLILISPAKTLDYQSPLTTTRYT
		1				LPELLDNSQQLIHEARKLTPPQI
						STLMRISDKLAGINAARFHDWQ
		ŀ				PDFTPANARQAILAFKGDVYTG
						LQAETFSEDDFDFAQQHLRMLS
				ŀ		/GLPYRLEMGIRLENA\RGKDLY
		1				QFWGDIITNKLNEALAAQGDN
		Ī				VVINLASDEYFKSVKPKKLNAE
						IIKPVFLDEKNGKFKIISFYAKK
						ARGLMSRFIIENRLTKPEQLTGF
						NSEGYFFDEDSSSNGELVFKRY
			<u></u>	<u> </u>		EQR
25421	55789	A	25562	3	320	IGLVLPISVRKRHAASARCLVN
1						RRPMPRLHGLPGIPMRPPSLPRV
						RHAGTVFRRHRKRGSPMAY*Y
						GHSDVVKHREALVADLCHKAA
			<u> </u>			FQLCRGQRHCRPALWRHR

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25422	55790	A	25563	1600	1908	
25423	55791	A	25564	142	326	TRRAPSAISSRCCCNSHQTSKSS LVRTSSSLRSSSHLRTIA*CSSV HLPISCMQIRPSQAR
25424	55792	A	25565	1	1599	
25425	55793	A	25566	393	611	HSRAVCRFYPAKTSPFPVRRCR YRQFVPR*HQNE*AGRE*QPCSI ATVLPYRALPSLAHPRLSADQL PAAHL
25426	55794	Α	25567	971	1628	
25427	55795	Α	25568	3	1895	
25428	55796	С	25569	1	2505	
25429	55797	Α	25570	1	3345	
25430	55798	A	25571	457	790	PSFRRRYRQVVVISVRKRHAAS ARCLVNRRPMPRLHGLPGIPMR PPSLPRVRHAGTVFRRHRKRGS PMAY*YGHSDVVKHREALVAD LCHKAAFQLCRGQRHCRPALW RHR
25431	55799	Α	25572	1662	1767	
25432	55800	A	25573	3	603	RLRSLFFLPDKMKLKDCVMIKK IFALPVIEQISPVLSRRKLDELDL IVVDHPQPPSGDYTLHPLPPIIR ANRPNRFRINAGCTRTFTLSTPI NVFRHKILRTQEFPSLTHHAYSP LLVVINKAKFDGLSPEFQQALV SSAQEAGNYQRKLVAED/LAK NHRRHERSGRGS/LSPISTAKPL ATHWGIRFATCLLKMCRRELIC
25433	55801	A	25574	2271	2519	HRQLVIVFQQPLVSGFHEAELS LDDAKWVLHP\ARMLAFMYSM LMAVLFLRGCCFKVLTLPGRSA ISQSTSTSASSSRCGAPW
25434	55802	Α	25575	931	1213	
25435	55803	A	25576		2767	MIAILSGLRCSIVANYIYAENNL FWLRLNNPKAIWCLRICTFATV IGGTLLSLPLMWQLADIIMACM AITNLTAILLLSPVVHTIASDYL RQRKLGVRPVFDPLRYPDIGRQ LSPDAPLTTTRYTLPELLDNSQ QLIHEARKLTPPQISTLMRISDK LAGINAARFHDWQPDFTPANA RQAILAFKGDVYTGLQAETFSE DDFDFAQQHLRMLSGLYGVLR PLDLMQPYRLEMGIRLENARG KDLY\QF\WGDII
25436	55804	Α	25577	395	467	
25437	55805	Α	25578	1	675	
25438	55806	Α	25579	1	4014	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ĺ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			1	sequence		
25439	55807	ĪA	25580	1802	2385	I IRKISAPALIKSRTG*EMASLPTS/
23439	33007		23300	1002	2505	HDKQIEQALADVDVPIVGVGGS
						YHLAESYPPVHYIATDNYALVE
						SAFLHLKEKGVNRFAFYGLPES
						SGKRWATEREYAFRQLVAEEK
						YRGVVYQGLETAPENWQHAQ
						NRLADWLQTLPPQTGIIAVTDA
						RARHILQVCEHLHIPVPEKLCVI
1						GVMNRRTVRSPFRESLKISP
25440	55808	A	25581	1	1568	MVKQLKLVSFLLRPGDSKQGY
23440	33808	I ^A	23361		1300	DKHSSLQVGLMIPVCSCIRFRKP
						SEQSFKFEEGTDAAFHGFWLW
		1				PCTMLNVTPIKGMMTVIYEFPV
						DSETTLQNYDIYFTNEELTDEQ
						KSLIEWYRDVFRPEDLRLVESV
						QKGLKSRGYRGQGRIMADSSG
	İ					SGISEHAWCMSRGAQGRQFCK
						GAKMSQVLITGATGLVGGHLL
						RMLINEPKVNAIAAPTRRPLGD
]					MPGVFNPHDPQLSDALAQVTD
1						PIDIVFCCLGTTRREAGSKEAFI
						HADYTLVVDTALTGRRLGAQH
						MLVVSAMGANAHSPFFYNRVK
						GEMEEALIAQNWPKLTIARPSM
						LLGDRSKQRMNETLFAP\LFRL
						LPGIILLEELNWKTWAERWNS
						MLGPRAVKVFFSLAFLPGTLEP
	ļ.					AQILFPTQRQGKKAALLLGLW
						KLEIYSDNRRSPEVSTAWNSPP
						TGKLDNKQYMIMIPRKRETNTT
						TRGASNARLDPPSRKGYKGHY
		1				ETIGHALEHDKQSQGSYNVEHT
	i					IKROTPRVRRCTSIIAGIYQELG
25441	55809	В	25582	1	1560	
25442	55810	A	25583	1	2115	
25443	55811	A	25584	3036	3957	RIYDAVSEPAEPVDCPRNPVCR
						FY*QYRDHRGKTVYDVASGDA
		į				LFISELG/PLPENVTWLSPEGEFQ
						KWN/GTAWVKDTEAEKLFRIRE
						AEETKNNLMQVASEHIAPLQD
						AADLEIATEEEISLLEAWKKYR
						VLNSRAMALILTPDLIRSNTACF
	}					CSSFVLAGQTAKAAADDGLFPP
						IFARVNKAGSPVAGLIIVGILMT
						IFQLSSISPNATKEFGLVSSVSVI
	-					FTLVPYLYTCAALLLLGHGHFG
	1					KARPAYLAVTTIAFLYCIWAVV
						GSGAKEVMWSFVTLMVITAMY
						ALNYNRLHKNPYPLDAPISKD
25444	55812	A	25585	1	2490	
	100012	1.,	,		1	l

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25445	55813	A	25586	368	761	LYCEACIMAFRMSEQPRTIKIY
25 (15	33013	` `	25500		, , ,	NLLAGTNEFIGEGDAYIPPHTGL
						PANSTDIAPPDIPAGFVAVFNSD
		l				ESSWHLVEDHRGKTVYDVASG
		l				DALFISELGPLPENVTWLSPEGE
		ŀ				FQKCSGTTRVKNAKA*TLF
25446	55814	A	25587	1	809	MPVSSYERGHAEKRVVQECCG
23440	33014	`	23307		1007	INLTSSANTILSDCERVNGYCVL
						CHTTRALVTGNHHRNAVYAET
						PSNSRINYIVSSHNAFRMSEQPR
						TIKVYNLLAGTNEFIGEGDAYIP
						1
						PHTGLPANSTDIAPPDIPAGFVA
						VFNSDESSWHLVEDHRGKTVY
						DVASGNALFISELGSLPENVTW
						LSPEGEFQKWNGTAWVKDTEA
		ŀ				EKLFRIREAEETKNNLMQVASE
		ŀ				HIAPL\QDAADLEIATEEEISLLE
						AWKKYRVLLNRVDTSTAQDIE
		ļ				WPALP
25447	55815	Α	25588	1	2301	
25448	55816	Α	25589	718	1131	
25449	55817	Α	25590	279	513	CRRIDTWQPIYQTCGRLMAVEL
						LTVVTHPLNPSQRLPPDRYFTEI
						TV/SHRMEVVKEQIDLLAQKAF
		ļ				SLAYLSPCESGL
25450	55818	Α	25591	1	407	PQDNAKKLRPHQLSSDEGHTG
						MGGTAKDVGFVSMRTYSWISS
						ALPKRLGGGVMPIGATIATEEV
		İ				FSVLLGNPFLHTTTFGGNPLAC
						AAALATINVLL\GRTYRLRLSK
1						KAICCWTVSVNWRGNIPIWYR
		<u> </u>				KRVVKGC
25451	55819	Α	25592	1	2079	
25452	55820	В	25593	1	700	
25453	55821	Α	25594	1427	1845	ACISGFSTETSASGRVSVITDAS
1						LFRVLPRVCGMAVAATKRSGRI
						TTSEALCCANQLRRRPCVSPTR
l						STPSWLSCAV*ICTSGTT*MKR
						VPNARHCWICSSLIPHSGLHPQP
						ITAKFAPIALSSSVTSPFSSVTFG *IN
25454	55822	A	25595	2	295	
25455	55823	A	25596	1	900	
25456	55824	A	25597	ti —	1422	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25457	55825	I	25500	10	535	VCNDEOD AVCCUTATIVI AID AL
23437	33823	Α	25598	2	333	YGNREQRAVSSHTNTWVNRAL ARGWSSLGYRWWVSAWTPSR
						RFISQLMSRPVDASAAKVKRSA
						SVPCGSIPFGNCFSVRLRILSASC
						GCIILPVRFSSSSGSEIPSTISSGSI
						TLPLDLDIFWPSSSRIRPVM*TV
						WNGTCGLPFSSLMKCMVIMIM
						RAIQKKMMSKPDTITLVGWN
25458	55826	Α	25599	442	1368	RRTMAAFACGVVRDASEVIRPS
						STLVPLVGEKHAKGLFGTIVDN
						FYLVALIFAMGTSLGLATPLVT
						ECMQWLFGIPHTLLGFI/LYNAI
						CVACGLQKGVRIASDVRSYLSF
						LMLGWVF\IVSGASFIMNYFTD
						SVGMLLMYLPRMLFYTDPIAK
						GGFPQGWTVFYWAWWVIYAI
						QMSIFLARISRGRTVRELCWIL
						WTVLGSNTLLLIDKNIINIPNLIE
						QYGVARAIIETWAALALNNGN
						MWGFFILWFIGPGTLVNAWQNI
						RAQFVHHILRRNGIADGFRHLA
						ALTIHGKAVGQYLTIRRFTLHR
25459	55827	Α	25600	2468	3907	
25460	55828	Α	25601	3	3738	YKTNYTDNSEYYANVGFNVKP
						VKRDGKMSVMSDPDTAIYPFPP
						KPTPLSIDEKAYYREKIKRLLTE
				•		RNAVMVAHYYTDPEIQQLAEE
				•		TGGCISDSLAMARFGAKDPASN
						LLVAGGGTCRSVVINAVFSYVT
						NVWGWAFEWYMVVMLFGWF
						WLVFGPYAKKRLGNEPPEFSTA
						SWIFMMFASCTSAAVLFWGSIE
						IYYYISTPPFGLEPNSTGAKELG
						LAYSLFHWGPLPWATYSFLSV
						AFAYFFFVRKMEVIRPS
25461	55829	A	25602	1331	1456	
25462	55830	A	25603	549	588	
25463	55831	Α	25604	19	1629	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ŀ		sequence		
25464	55832	Α	25605	1	1995	MVGVRFALGLMWLTLIVAETIS
						ANSGIGYLAMNAREFLQTDVV
						VVAIILYALLGKLADVSAQLLE
						RLWLRWNPAYHLKEATLSAEQ
		ŀ				AARIREMTVITSSLMMSLTVDE
						SDLSVHLVGRKINKREWAGNA
		ĺ			i	SAWHDTPAVARDLSHGLSFAE
		ŀ				QVVSEAHSAIVILDSRGNIQRFN
						RLCEDYTGLKEHDVIGQSVFKL
						FMSRREAAASRRNNRVFFRSGN
		ļ				AYEVENVPDPGVEQADDIILRIT
						ATAICGSDLHLYRGKIPQVKHG
						DIFGHEFMGEVVETGKDVKNL
						QKGDRVVIPFVIACGDCFFCRL
						QQYAACENTNAGKGAALNKK
						QIPAPAALFGYSHLYGGVPGGQ
						AEYVRVLKGMWGRLNDAQPG
						FRSSRPLRYGAGIMTLLNISGLS
		l				HHYAHGGFNGKHQHQAVLNN
						VSLTLKSGETVALLGRSGCGKS
	1					TLARLLVGLESPAQGNISWRGE
						PLAKLNRAQRKAFRRDIQMVF
						QDSIST\VNPQPNRLHIV/RFPLT
						RLNQYQPSFSTDLYHGTLKPAS
						LTMSGVVLLGRM/PRKTSQQV
						QWA\CIARTDKASMPTVYSESL
						FVEPLERVLYVAVILTQIIRGLR
				:		DTEIETINPDIAVHFALCFHGLG
						SIPTKGAEKSPFLTIHKVRVKPD
						IPSPVSIMAPCRAHPTCEALFNN
						STALHSPWFCDSL
25465	55833	A	25606	179	371	CVLQYTHWPACSTVA\HSRRW*
						NRRRLRLAVTKTGSLFSHRCAG
						ARLAFSKKLRCRDALPASRS
25466	55834	A	25607	544	983	SVQNPRVNWIHAALQRTGRGR
						RRHEQHGEDHFVNGAAGVHQ
						AANGLVNPPRHQVFGAHQAKG
						DGENHRQRGAPDGDLQRDGHF
						GEVILPLAEIGREEVGGERRHV
· ·						AAVFDQ/S*AGPFPRPATRRPTR
						RVQRPSSEARTSCAWVGRW
25467	55835	В	25608	1	1746	(
25468	55836	A	25609	1	858	-
25469	55837	A	25610	98	288	LCRNNGEPLARSGAAMGALLD
]						REHSRHP\PDLEG\GSQQQVCDS
						GGSVTAANTSGGQQVKAQLEL
L	<u> </u>	L		L	L	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25470	55838	IA	25611	11934	2812	RPQTYSDAQPGFRSSRPLRYGA
						GIMTLLNISGLSHHYAHGGFNG
						KHQHQAGLNNVSLTLKSGETV
						ALLGRSGCGKSTLSRVLVGLES
1						PAQGNISWRGEPLAKLNRAQR
						KAFRRDIQMVFQDSISAVNPRK
						TVREILREPMRHLLSLKKSEQL
	Ì					ARASEMLKAVDLDDSVLDKRP
						PQLSGGQLQRVCLARALAVEP
<u> </u>						KLLILDEAVSNLDLVLQAGVIR
						LLKKLQQQFGT\ASLFITHDLRL
						VERFCQRVMVMDNGQIVETQV
						VGEKLTFSSDAGRVLQNAVLP
	1					AFPVRRRTTEKV
25471	55839	A	25612	1	515	MLAVECRSLPFPPLVQPVRYTD
23471	33637	' '	23012	1		VTVKDDYNPETEQYTLTISQRT
	1			İ		PATPDQAEKQPLHIPFAIELYDN
						EGKVIPLQKGGHPVNSVLNVTQ
						AEQTFVFDNVYFQPVPALLCEF
						SAPVKLEYKWSDQQLTFLMRH
						ARNDFSRWDAAQSLLA/ILHQA
						ERRASSARSAAVSAGACG
25472	55840	A	25613	1	1053	
25473	55841	A	25614	1	1521	
25474	55842	Α	25615	430	1120	QTASTFLKRGKCASRQSKLCGN
						RRRKRRQRISSYCLLRGTACSP
						AHQPDAFHQKRGNAGRHYAEC
		1				VLQYSSCSTRYS\PGR\TQLINLF
	1					EVADGKRLVDLPGYGYAEVPE
						EMKRKWQRALGEYLEKRQSLQ
						GLVVLMDIRHPLKDLDQQMIE
						WAVDSNIAVLVLLTKADKLAS
						GARKAQLNMVREAVLAFNGD
						VQVETFSSLKKQGVDKLRQKL
						DTWFSEMQPVEETQDGE
25475	55843	Α	25616	407	1176	AYSTSLRVKICRKTSTLLLRSRV
						TQIRSNTKSTKRAAHCSLTASC
						PPRCSIHATTVTSTTPCLWRRIA
						TMLRALARLLLRICFSGRTLKIA
						CLLLLVAGATILIADR/AALYHA
						GKVKWLLVSGDNGRKNYDEA
						SGMQQALIAKGVPAKVIFCDY
						AGFSTLDSVVRAKKVFGENHIT
				1		IISQEFHNQRAIWLAKQYGIDAI
	1					GFNAPDLNMKHGFYTQLREKL
						VIAKGHGSLLTDLPVAKAGTM
	<u> </u>	<u> </u>				WAIEIDDVARHSRSHL
25476	55844	A	25617	3	168	NPFYSAMR\KSAGIRHILARHVE
						GASHMAEGYTRATAGNIGVCL
						GTSGRCCFRLCW
25477	55845	В	25618	169	1227	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence	:	07/340,217	sequence	peptide sequence	detection, 1 possible nucleonide insertion)
25478	55846	Α	25619	489	740	AIARSCADYHPANFRHHRAAA
1			}			RAGDDYLSRRAERQPGAKAGG
ĺ						SRLRAGKRPCSAFRYW*CAAG
						E*SVGEKIMAGDVISPKRRL
25479	55847	A	25620	146	475	
25480	55848	В	25621	38	191	
25481	55849	A	25622	662	4568	
25482	55850	Α	25623	20	2661	AGHIHRPRYQHGVGTESTSIMS
						VASRSAERVFLTKFESWKNFPI
		1				NRGRNKMAKMRAVDAAMYV
		İ				LKKEGI\TPAFG\VPGAA\INPFY
						SPMRKHG\GIRHILARHVEGAS
						HMAEGYTRATAGNIGVCLGTS
						GPAGTDMITALYSASADSIPILC
1						ITGQAPRARLHKEDFQAVDIEAI
		ļ			1	AKPVSKMAVTVREAALVPRVL
						QQAFHLMRSGRPGPVLVDLPF
						DVQVAEIEFDPDMYEPLPVYKP
						AASRMQIEKAVEMLIQAERPVI
						VAGGGVINADAAALLQQFAEL
						TSVPVIPTLMGWGCIPDDHELM
		ł				AGMVGLQTGHRYEKYTEGRKI
						VHIDIEPTQIGRVLCPDLGIVSD
1						AKAALTLLVEVAQEMQKAGRL
						PCRKEWVADCQQRKRTLLRKT
	Ī					HFDNVPVKPQRVYEEMNKAFG
						RDVCYVTTIGLSQIAAAQMLHV
						FKDRHWINCGQAGPLGWTIPA ALGVCAADPKRNVVAISGDFD
						FQFLIEELAVGAQFNIPYIHVLV
1						NRPAANGAQGSCHPDAPAVFR
						SSIFIRKSEEKLVVTMNELSASR
						ALPEETMPGRFELKPTLEKVLH
1		1				APDNFLFMDPLPPMHRRGIIIAA
						IVLAVGFLLPSDDTPNAPVVTR
		1				EAQLDIQSQSQPPTEEQLRAQL
		1				VTPQNDPDQNLYRKASQKSSR
						KLPKRDLPHILTPFRAPMASVC
						AKRKGQVLHQQYSLKKYRYK
25483	55851		25624	1	570	ARIGOGVENOGISERRIKI
25484	55852	A A	25625	312	873	APRSRVQAKIAASNTGELNALQ
23404	22022		23023	312	073	QLGFS\LVEGEVDLAL\PVNNA\
						SDSGAVVA\QETDIPA*RQLASA
						AFAQSRFRAPWYAPDASSRFYA
						QWIENAVRGTFDHQCLILRAAS
						GDIRGYVSLRELNATDARIGLL
						AGRGAGAELMQTALNWAYAR
						GKTTLRVATQMGNTAALKRYI
						QSGANVESTAYWLYR
<u> </u>		<u> </u>		<u> </u>	<u> </u>	QOUANVESTATWLIK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25485	55853	Α	25626	2	922	FSPPGLHDAGEAVFWRVNKSA
ļ						NYGVLDKLARGYADLSKAESQ
						WDEMMRTAGSLKLGTIHASINS
						LFENITAAPADPILGLADLFRAD
1				İ		ERPGKINLGIGENPSCCRTSSSP
1						VFDAAIWPAPVNNASDSGAVV
						AQETDIPALRQLASAAFAQSRF
						RAPWYAPDASSRFYAQWIENA
						VRGTFDHQCLILRAASGDIRGY
			1			VSLRELNATDARIGLLAGRGAG
						AELMQTALNWAYARGDHMI/R
						I*RTAGGGNRTRLYAVGNG*RQ
						TVWRWRFYPSLPAVAGATFWQ
						RQSVTDAVLHRFAGDGGAAAR
						YPAWR
25486	55854	A	25627	911	1441	AIPSISCFLLAIKALILSAIWLNA
23400	33034	`	23027			TPSRSKLEPSSKWIRSLRCPSPN
		1				RCAAASSFNISCQWFLSEYGPP
		1				NQVASGKTSPFVIVFKQPRYRA
						VTKPNCRLMMRNGCSTLARML
						PVMYAMLMPVLFLRGCCFKVL
		1				TLPGRSAISQSTSTSASSSRCGA
						PW*PASAETNCSSPCSRLPS
25487	55855	A	25628	1	1247	MERVLDNLLNNALRYCHSTVE
23407	33633		25020	1	12-17	TSLLLSGNRATLIVEDDGPGIAP
						ENREHIFEPFVRLDPSRDRSTGG
				ŀ		CGLGLAIVHSIALAMGVNAQA
]			DFMKLTGADEQIAAYCFKMPF
		ŀ		ļ		VFHLIEEVSRNFGYALCLINIDH
						KAALNITQRTAANILLDGQMQ
						HAVDQAFTQRTARQGSTFVQY
						ARLARKLATANICWSYFVPVR
						ASIEPLTWENAFFGVNSAIVRIT
						SEAPLLTPDALAPWSRVQAKIA
						ASNTGELDALQQLGFSLVEGEV
İ			ļ			DLALPVNNASDSG\AVVAQETD
						IPALRQLASAAFAQSRFRAPWY
						APDASSRFYAQWIENAVRGTFD
			}	ŀ	ļ.	HQCLILRAASGDIRGYVSLREL
			1	1		NATDARIGLLAGRGAGAELMQ
			1			TALNWAYARGKTTLRVATQM
				1		GYTAALKRYIQSGANVESTAY
						WLYR
25488	55856	A	25629	1	1925	WLIK
25489	55857	A	25630	402	664	TARRLLFFRD/RTARVAVPNAA
23409	33031	^	23030	1702	1004	VVLPFPLPVKTMIIPRSLGRRRS
				1		GGICRPLGYLVAMDAIDGLQN
						ADNSNGESRIATSSFRSRSARR
L	L	Щ.	1		<u> </u>	ADIOTOLORIA 1001 ROROARK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25490	55858	A	25631	920	1237	IARPGCHLSHCHSEPSGRLSK*P
						FSCNSLKTGDF*PI*GLIQTQIFS
		1				PFAFRRIIPSGSGKVSGSHSKS
		1				HHWKAFIQKQSKWKTCSGRLR
		1				SAIPSIKLFTVASS
25491	55859	Α	25632	1	1341	MVHRFGIRVRFVALGFYQDVIL
		ļ			·	PVRAHRHEWGHYVHLCASIQR
	:					QIDVSFARDATSHSSCPVTGVR
						FWGYPFAANKVVIAGFELEGET
1			}			MTRPGLICAEGGLLVMIVSLRS
			ļ			ELYDSYHEFTRQRVTEIVEKAA
						NHLPAVVEMMTLHRCAINRHN
		1				FVRRVGPAGISTFFASSPDGVN
	1					AGFSTFRVVFTCQRSRPLQRHS
						HADVQTATLLLNTVNLEQKQA
						NAILSGLSDMIPNSSPESAPEIQL
		İ				LQSRMILGKTIAELNLRDIVEQK
						YFPIVGRGWARLTKEKPGELAI
						SWMHIP*TLSKNRPTPFS/DGLS
						DMIPNSSPESAPEIQLLQSRMIL
	1					GKTIAELNLRDIVEQKYFPIVGR
1						GWARLTKEKPGELAISWMHIP
		ļ				QLNGQDQQLTLTVGENGHYTL
						EGEEFTVNGMVGQRLEKDGVA
						LTIADIKAKPGTQFVLSQRTELE
						AINALQETFTVSERSKESGMLE
ļ						LTMTGDDPQLITRILNSIANNYL
						QQNIARQAAQDSQSLEFLQRQL
		İ				PEVRSELDQRKKNSTFIASSAIR
25492	55860	Α	25633	170	935	LTQEDTRINLIPMAVESLHLPTQ
		İ				MLINGGETYAWRNWKRGGQY
						KSATLNTGQPHVRARELVDSD
						TLASDAARLTCRHGLALPALFT
						LIAVVTSLLSPQFIRQILWSRLT
						KTR*AGDQLDAYSTTEWSGSA
						TDTHGWGKRPLYTGR*RVHRQ
						WYGRPASGKRWRCADYRGH*
						GQTRNTVCPEPAYRTGSD
25493	55861	В	25634	42	755	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ĺ		sequence		
25494	55862	A	25635	793	2127	RYCKGTVMDQIRLTHLRQLEA
23494	33802	ΙΑ .	23033	193	2127	ESIHIIREVAAEYSNPVMLYSIG
		1				
						KDSSVMLHLARKAYYPGTLPFP
				•		LLHVDTGWKFREMYEFRDRTA
İ						KAYGCELLVHKNPEGVAMGIN
						PFVHGSAKHTDIMKTEGLKQA
						LNK/YTTSRAKERIYSFRDRFHR
						WDPKNQRPELWHNYNGQIN\K
						GESIRVFPLSNWTEQDIWQYIW
						LENIDIVPLYLAAERPVLERDG
						MLMMIDDNRIDLQPGEVIKKR
		1				MVRFRTLGCWPLTGAVESNAQ
		Ì				TLPEIIEEMLVSTTSERQGRVID
						RDQAGSMELKKRQLIELIINVH
						NLLKNGFGFQVKVNRIALLAIN
				-		VEFFFRLVQLAAHFSSGTLAQD
						KLCSWFGLNVRDSQRNAIFFQT
						LADHTIDGELFTFQCIVAVFPNL
		İ				PIYSYATIYVSSSSIPISSEESPSIR
						FACSQTSRQPPVSDSRFCVSQV
25495	55863	A	25636	360	566	GAETMGDLLWVVGILLMCSLS
						TLVLVWLDPRLKS*RT*A*CGA
						LTSHQAINNGRNEKRKPIRESDP
						LKA
25496	55864	A	25637	1	312	LLRKSVSVIIKGTIKTIIGFMLLQ
						AGSGILTSTFKPVVAKMSEVYG
ĺ						INGAISDTYVMQDG*APELGLS
						PGRFLRHDTDFVLQLGIVMVRL
					:	AGDGSLNINAQANN
25497	55865	A	25638	517	796	PTARFGWVKGVDLPAD/CNRQ
23471	33603	``	25050	317	1,70	DRFAIDVRVQGS*\WGCVLPLF
	İ					RVLLFVSCALLLVSVSCVLRLL
						FCCSS\LSLFLSGLGGGRSPKAG
	1					DVFRDRR
25498	55866	A	25639	47	411	AIKVRELSIKLKMC/YSLIKTLH
23490	33800	^	23039	147		G*KYRNFMVIGSMVFGKKAITH
1						VAGVKRRTAVVVGFKVFDIQR
						SQFRIVVTGDLAQLLYRVIKVIT
1						CGHFVRQHGIVLCAGVLHVGD
						RYQAHVKTLGGLI
25400	55067	ļ.—	05640		1400	RYQAHVKILOGLI
25499	55867	A	25640	1	1488	CCICL/VSVADIDVIV/VIC/WATTA
25500	55868	Α	25641	$ ^2$	298	CCICL/YSYAPIDVW/WG/WNTA
						GQVQAA*TLKAASSSIVTIPLRN
						TFIFNYSLSGLRRRRLIIRSRRHS
						THASNITLIAASTASRIGWFASA
0.555	55060		25645	ļ,	2064	YHTRSNHGR
25501	55869	C	25642	1	2064	
25502	55870	В	25643	1	1116	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25503	55871	A	25644	312	620	SACRCCLVVVPLFVCL/GSSLSL
23303	338/1	^	23044	312	020	APTFPLLASFSSSRASRVVAFGL
		ŀ				GSLRAGSALCLPCAAVVLALRS
						AGTCLEWWWWHSVVLPPETK
						RLPSTSLIPYPASVSI
05504	55070	!	25645		2227	RLPS15LIP 1PASVSI
25504	55872	A	25645	2	2237	DA CONCOUNT OVER CONTACT IN DATE
25505	55873	Α	25646	692	6383	RMSRSSDYDQYRSRNALIRRHI
		ŀ				EKMDASLHVGTKEFDISKVSEV
	1					DSVDDLLIDNAARYLLKDWKG
						VGELVNGVEVALEYTAERGIAL
1	-					LKQNPELYWQILAEAASIAQVG
		l				GEVWIAPKFERNGLRLLANSEI
		ł				GAASQLQTYHRVDPGRPCKLD
						LSGGMSQRVMIAMAIACRPKL
						LIADEPTTALDVTIQAQIIELLLE
						LQQKENMALVLITHDLALVAE
				1		AAHKIIVMYAGQVVETGDAHA
						IFHAPRHPYTQALLRA
25506	55874	Α	25647	856	1467	
25507	55875	Α	25648	1823	2014	SQEIALMQFRPRLTGASGSP*RP
						PHRTLLRSDQNAATIPQKRASA
		ŀ				LIPLMRILPVRYGHLLLC
25508	55876	Α	25649	1	218	
25509	55877	Α	25650	2740	3293	IFAVNLILQCHVGGTHHQRFLL
						FAGNGDAGNQIRERFPNAGG/E
						LQSPDVALLLRPAFWRRPQSSA
						AAAHGE*SRGSVPEAPYTRRQF
						ALPAPPIKCAPLPERQRPSVPSR
						HLRLPACPGYKNRASSAKPSYS
						FLAQRQHLQRLSPLPQNQNLPQ
						RRRDQRQRAGAMSARVPDLISS
						SRSMRRTS
25510	55878	В	25651	1	1875	
25511	55879	В	25652	1	2981	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25512	55880	Α	25653	11	1649	MTFRNCVAVDLGASSGRVMLA
		1				RYERECRSLTLREIHRFNNGLH
						SQNGYVTWDVDSLESAIRLGL
1						NKVCEEGIRIDSIGIDTWGVDFV
						LLDQQGQRVGLPVAYRDSRTN
1						GLMAQAQQLGKRDIYQRSGI
						QFLPFNTLYQLRALTEQQPELIP
						HIAHALLMPDYFSYRLTGKMN
		į				WEYTNATTTQLVNINSDDWDE
						SLLAWSGGQQSLVWVAEASGE
-						WGVLACTLFAWNRNYFVLLFV
						GVFLSSFGSTANPQMFALAREH
						ADKTGREAVMFSSFLRAQVSL
						AWVIGPPLAYALAMGFSFTVM
						YLSAAVAFIVCGVMVWLFLPS
						MRKELPLATGTIEAPRRNRRDT
						LLLFVICTLMWGSNSLYIINMPL
						FIINELHLPEKLAGVMMGTAAG
						LEIPTMLIAGYFRQTSGPTAYGR
						SVGRIRRSRRIRQSCYYQRIFSL
						LACELANALERPENFCGMHFFN
						PVHRMPLVEIIRGEKSSDETIAK
1						VVAWASKMGKTPIVVNDCPGF
						FVNRVLFPYFAGFSQLLR/ERRG
						FPQDRQSDGKTVWLADGPGIS
						AGRCGH
25513	55881	Α	25654	1590	2855	RNLLGEAADPVSGPHSAIYRPD
						WPKSGDCACQGPIKGSANRPA
						VVPHVHGP\SEDLEVFLNVFGE
						LPQPLIDTQILAAFCGRPMSWG
						FASMVEEYSGVTLDKSESRTD
			!			WLARP\LPERQWEYASPNVWY
						LLPITAKLMVETEASGWLPAAL
						DECRLMQMRRQEVVAPEDAW
						RDITNAWQLRTRQLACLQLLA
						DWRLRKARERDLAVNFVVREE
						HLWSVARYMPGSLGELDSLGL
						SGSEIRFHGKTLLALVEKAQTL
						PEDALPQPMLNLMDMPGYRKA
1						FKAIKSLITDVSETHKISAELLAS
						RRQINQLLNWHWKLKPQNNLP
						ELISGWRGISAVKSSEAGLGAL
						SATSGFGKLQIQHQAFTLRTRT
						TDQHTLLAGTLFIAKILIPMVSF
						TFQYCCFAVPQATYCHRIAPIL
	<u> </u>	l	l		<u> </u>	GNGAMKPGRKIS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25514	55882	A	25655	1	3315	MDELRLVLMTIIHIKLPPIRRAD
						GSFGNDLTTGAGFNDRKTVPR
]		YARVHVYRCRKVICIFTKRIPFC
		İ				LDVPAHRALVVMKGTLYGMP
						VVAAAFEFAFMGGSMGSVVGA
						RFVRAVEQALEDNCPLICFSAS
						GGARMQEALMSLMQMAKTSA
						ALAKMQERGLPYISVLTDPTM
						GGVSASFAMLGDLNIAEPKALI
						GFAGPRVIEQTVREKLPPGFQR
		ŀ				SEFLIEKGAIDMIVRRPEMRLKL
		ŀ				ASILAKLMNLPAPNPEA
25515	55883	Α	25656	246	422	RHADTDLRR*ISDRFSPAPPADP
						ANGGR*TPRHAAKRRDPGTPV
						QPARKPSKPTRLLR
25516	55884	Α	25657	884	1276	LVTRQLSPGNRCNVFAFSSAVE
		1				TSLTCQGWLSPVCAVCQAISSS
		l				HNASIQVRSPAASCGSPFCSKRP
		i i				SALWWTSVSPFLSPRRRTSGAN
		l				AESVSAT*SKRRSSARRSASARS
						TLSGAALRSRTASNEWRS
25517	55885	Α	25658	649	1427	LPLLGWSMSPRMTAQLACDAL
		1				QMALWRRKRPRNVIVHTDRGG
		l				QSVQQFRALKRHITASRETERF
						YLDVQKQFPKVTAQKVIVSEA
						GASVYSASELAAQEFPDLDVSL
		1				RGAVSIARRLQDPLAELVKIDP
		ļ				KSIGVGQYQHDVSQTQLARKL
						DAVVEDCVNAVGVDLNTASVP
						LLTRVAGLTRMMAQNIVAWR
						DENGQFQNRQQLLKVSRLGPK
						AFEQCAGFLRINHGDNPLDAST
						VHPEAYPVVERILAATQQALKD
						LMGNSSELRNLKASFLDNEEN
						WPLLLDALIPVANTCEMILMPA
						CFGLADDKLWRWLNEKLPCSL
						MLLPTLPPSVLGIRLQNQLQRQ
						FVRQGGVWIECGDGGDTTTITI
						TQR**PAGRVYRSPGSLSGGGT
		<u> </u>				HSGSNTAGTERSDG
	55886	Α	25659	1	1878	
25519	55887	Α	25660	1	4329	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ 1D NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25520	55888	A	25661	2514	4173	PTSCLPDDYVGHCRGVSAAHF AGSVSCVRLLSACARPLWRHL GLALLRVWCRTFLLLSALASVF CPCSVVWGRLLFSASLSVVVW GGVFLSPLGSLSPRGTARLGAA CRSRED/RRR*WRGIFNRDTFAS AVPASVNPDKPCAPAACIRFTS TSAY*VGCSDRNA/DAEASGEG WRRFGDATLSTGQFGGEAREE VILGLIGSQA\KTGGNTPKASAV RKITLVAWPALETGLTMLSM** IG*ETRVFSVFEPSSKSIVPSSRT VTFSSNASRRIAL*ISGSASFDSL IVFA*QPPSKLNTPSSSQPCSSSP IRRRFGSVERVVLPVPERPKNT ATSPFSPTLAEQCIEAIPSAVFA APVRVAASMIISGFCALASIRPS ASTRRPSASVFITSTFLPLR*VMI SPSLNALPLIRLSAQQRNSFTRL FSPRVMANASAPVTVAAPPMS DFIESINAVCLMQARVHWRFFI ASWVVIQPPAIPENVLNRVRLC AAPGTLCLTTFVSSSGDR*NAL CNRCNRPAPCFRSSAIRFATVLA ARRHDWVWPISPLIPRPSS

NO: of peptide sequence load in USSN. voice first conduction of first conduction of peptide sequence addon for last amino acid decision, "possible nucleotide insertion) acid methods in section." 23521 55889 A 25662 2221 4400 RGCVVSGKSSCSKTNLPSDDKF TGASCPLRDLE FRYRAHALY. 23521 55889 A 25662 2221 4400 RGCVVSGKSSCSKTNLPSDDKF TGASCPLRDLE FRYRAHALY. 23521 23521 4400 RGCVVSGKSSCSKTNLPSDDKF TGASCPLRDLE FRYRAHALY. RELEARCH TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGASCPL TGA	SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence	NO:				1	1	
25521 55889 A 25662 2221 4400 RGCVVSGKSSCSKTNLPSDDKT TGASCPLEDLPLRYLRAHALVI AEQLAHEFSLGIAIVEEQLQQLI EQGLVMLQQDIWVSDEWER LRLRSLQAAREATRPVAATTY/ RLLLERQGVVPATOGPOWNERS LRLRSLQAAREATRPVAATTY/ RLLLERQGVVPATOGPOWNERS LRLRSLQAAREATRPVAATTY/ RLLLERQGVVPATOGPOWNERS LRLRSLQAAREATRPVAATTY/ RLLLERQGVVPATOGPOWNERS LRLRSLQAAREATRPVAATTY/ RLLLERQGVVPATOGPOWNERS LRLSVASTWAS SAINASL WESOGPEVYGKPVYA GENYSQGVYYLASFANKIWLS. QGVVDLHGFATTNGLYYKSALDE KLLVYTAANRQIPAEQVFPGA QGLLEGLTKTGGDTAKYALDE KLLVALASASERKALTKEFOW SKTDKNYRAISYYDYALKTPA DTGDSIGVVFANGAIMDGETT GNVGGDTTAAQIRDARLDFKV KAIVLAVNSPGGSVTASEVIRA ELAAARAAGKPVVVSMGAA RSGGFGVITTVENSAATER GNVGGDTTAAQIRDARLDFKV KAIVLAVNSPGGSVTASEVIRA ELAAARAAGKPVVVSMGAAA ASGGYWISTPANYIVANPSTLT GSIGFGVITTVENSATER STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAADARI STPEOJIDKIAQGHVETAAAAADARI STPEOJIDKIAQGHVETAAAAADARI STPEOJIDKIAQGHVETAAAAAADARI STPEOJIDKIAQGAHVETAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA		sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
TGASCPLRDLELRYLRAHAL VI AEQLAHEFSLIGIATIVEQLOQUIL EQCLVMNLQQDIWYSDEVFRR LRLRSLQAAREATRPVAATTY/ RLLLERQGVLPATDGSPALFAS TSRGYYEGVDGVMRVIEQLAA SAINASLWESQEFVYGKPYYAN GENYSGGYYLASFANKUS QGVVDLHGFATNGLYYKSAUPFI RDDMSPAAREADSRWIGEL KLEVSTHYRTGTYKSAVEFFI RDDMSPAAREADSRWIGEL RUTVANARQIPAEQVFPGA QGLLEGLTKTGGDTAKYALEN KLVDALASSAEIEKALTKEFGW SKTDKNYRAISYYDYJALKTEA DTGDSIGVYFANGAIMDEG GNVGGDTTAAQIRDARLDPKV KAIVLRVNSPGGSYTASEVIRA ELAAARAAGKPYVVSMGMA ASGGYWISTPANYIVANPSTLT GSIGIFGVITTVENSLDSIGVHTI GVSTSPLADVSITRALPPFAQI. MMQLSIENGYKRFITLVADARI STPEQIDELAGAGHVAVTGGD DDAVAKAAELAKVKQWHLEY YVDEPTFFDKVMDNMSGSVRA MLPDARQAMLPAPLASYQVRS PCLTYILLLERYHSVQYHAKEN LRCLHGRDHRDAAFRAGLYTG VRSSTTPTGADAGIPSGODARF AIHEYIPRNGSSRFNPGKLAPH/ 25522 55890 A 25663 I 1665 25522 55891 A 25666 1761 1970 25523 55893 A 25666 1761 1970 25524 55894 A 25667 455 1137 GTKPRIDHILGTFTTGMPKG KLDVYAAPELPLKLLGRTTEG YRESFISLPVVNYLKDKLISNP V/RLDGYDVAGGELV/VLENVR FINGGKKDDEHRAQASTHGIC K/FADVACAGPLLA/AUDSLS //ADQLIVGGGIANTFI/AQGHL YGKSLYEADL/OFAKRLLTC NIPVPSD/VRVATEFSETAPATL KSVRHSDKISYISTGGGAFLEL WKYKYLPAVAMLEERG 25527 55895 A 25669 1496 1631 25528 55896 A 25669 1496 1631 25528 55896 A 25669 1496 1631 25522 55897 A 25667 462 1152					sequence		
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PCLTYILLERYHSYQYHAKEIN						,	YVDEPTFFDKVMDNMSGSVRA
LRCLHGRDHRDAAFRAGLYTG							MLPDAFQAMLPAPLASYQVRS
LRCLHGRDHRDAAFRAGLYTG							PCLTYILLLFRYHSYQYHAKEIN
VRSSTTPTGADAGIPSPGDARF							LRCLHGRDHRDAAFRAGLYTG
25522 55890							
25523 55891							AIHEYIPRNGSSRFNPGKLAPHA
25524 55892 A 25665 242 3059	25522	55890	A	25663	1	1665	
25525 55893	25523	55891	Α	25664	475	1182	
25526 55894 A 25667 455 1137 GTKPRDIDHILGTFITPGMPKGC KLDVYAAPELPLKLLGRPTEGE YNEEF/SLLPVVNYLKDKLSNP V/RLDGVDVAEGEL/VVLENVR FNKGEKKDDE/HRAQASTHGIG K/FADVACAGPLLA/AVLDSLSK /IADQLIVGGGIANTFI/AAQGHE VGKSLYEADLV/DEAKRLLTTC NIPVPSD/VRVATEFSETAPATL KSVRHSDKISYISTGGGAFLEL WKVKYLPAVAMLEERG 25527 55895 A 25668 1108 1932 25528 55896 A 25669 1496 1631 25529 55897 A 25670 462 1152	25524	55892	Α	25665	242	3059	
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K/FADVACAGPLLA/AVLDSLSK							V/RLDGVDVAEGEL/VVLENVR
							FNKGEKKDDE/HRAQASTHGIG
							K/FADVACAGPLLA/AVLDSLSK
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25529 55897 A 25670 462 1152			_			<u></u>	
		-	-				
L.,	25530	55898	Α	25671	545	598	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the cont	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
	<u> </u>			<u> </u>		
25531	55899	Α	25672	1	1761	MRGHNWIDMDQARQQSTPHT
						DRKICHKNDRPRTTPLRKIDAH
						PHTLIILPAESTLSTTKYKLRAL
						LVENTNKDKSAVSVVFRGSVL
ĺ						HQQAANFPNLMLCFQLKNGVN
ĺ						ACNDFIMLFVDDNAAGSIERAA
ĺ						SPCINEKQHAFSESVPAIAAVRG
ĺ			•			LGSMIAVEFNDPQTGEPSAAIA
ĺ						QKIQQRALAQGLLLLTCGAYG
						NACKRRIDYVCRIKHSRRIRQV
						VLLNFAKSGAFSTTRGTDDKLT
						PVIQPRASGEYNEEFSLLPVVN
						YLKDKLSNPVRLVKDYLDGVD
1						VAEGELVVLENVRFNKGEKKD
						DETLSKKYAALCDVFVMDAFG
[TAHLRAGFLLTGI\GKLRCSLRC
ĺ		1				AGPLLAAELDALGKALKEPARP
İ					i	MVAIVGGSKVSTKLTVLDSLSK
1						IADQLIVGGGIANTFIAAQGHD
						VGKSLYEADLVDEAKRLLTTC
		İ				NIPVPSDVRVATEFSETAPANL
						KSVNDVKADEQILDIVRDLDAA
-						NVVINAVFSYVTNVWGWAFE
						WYMVVMLFGWFWLVFGPYAK
						KRLGNEPPEFSTASWIFMMFAL
		1				LYVCCRTVRLGREIHRRAQIVH
		l				IDIEPTQIGRVLCRISVLSLMLKR
25532	55900	A	25673	1	984	
25533	55901	A	25674	1	1181	MMAVSAPYVGDNIDDGGIYTQ
						SETHTISSDCVLDTAAMWTPLS
			}	i		AVIGCTTNPAQYPANAAEVYN
		1		:		KDGNKLDLYGKVDGLHYFSDN
						KDVDGDQTYMRLGFKGETQVT
		l				DQLTGYGQWEYQIQGNSAENE
<u> </u>						NNSWTRVAFAGLKFQDVGSFD
						YGRNYGVVYDVTSWTDVLPEF
						GGDTYGSDNFMQQRGNGFATY
						RNTDFFGLVDGLNFAVQYQGK
						NGNPSGESFTSGVTNNGRDALR
						QNGDGVGGSITYDYEGFGIG\Y
						TQTYNATRVGSLGWANKAQNF
]		EAVAQYQFDFGLRPSLAYLQSK
						GKNLGRGYDDEDILQICCMSDP
]		TQNRQASAKLALPCQVALLSSA
				1		SPHHAFTPSTSIIQPAHIIIHSSPP
]						PWASSHSISTHQQHTQCSSSSNT
1						
<u> </u>				<u> </u>		LAVH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	1 '	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25535	55903	A	25676	683	947	 QVRDSLQQGRERRIALSIGWLV
	00000					WREGCQDQCELEAGGASALTR
						RGSCRELDPAPFATMLLFCPGC
						GNGLIVEDG\QRCHASTDAVPP
25536	55904	A	25677	12	393	RGSCRELEPAEFETMLLFCPGC
						GNGLIVEEGQRCH/RAVACNT\
	-					CPYVHNITPQGNTIWKYPKLKE
						CGSDVLGGAAAW\ENVDSTAE
						SC\PKCEHPRAYFMQLQTRSAD
						EPMTTFYKCC\NAQCGHRWRD
25537	55905	Ā	25678	234	567	YVDETDDCFENNRCGQPVNGA
						YSPAGLSRVAADASLTTPHAK*
						TLRSLTLHQHDKTRTELLNDVA
						GALALDDKLGRSTNQLSGGEW
İ						TPSLARPFLLLGPRKMRLLPLH
						LPP
25538	55906	Α	25679	1	794	HCDGKHFTGVGKIMSIGMQLQ
						DVAESTRLGPLSGEVRAGEILH
						LVGPNGAGKESLTGREWPEMP
						TVKGTIQFPGHPLEAWSAKK\L
						ALLRAYL\SQQQTPPFPPPVWH
						YLTLHQPDKTRPELLNDVAGA
1						LALDDKLGRSTNQLSGGEWQR
						VRLAAVVLQITPQANPAGQLLL
						LDEPMNSLDVAQQSALDKILSA
						LCQQGLAIVMSSHDLNHTLRH
				ŀ		AHRAWLLKGGKMLASGRREE
						VLTPPNLAQAYGMNFRRLDIEG
						HRMLISTI
25539	55907	Α	25680	12	414	RAAPSRNLPPRMPIPIAAPIAAR
						PIITAAAMYSRSIFRFSMTVSSLF
						QLKRSSVGKLMFFRRHRQIDDR
						QNHEDEGLQRNDQDVENGPRH
						IQDPL*PPRQETSNQNEDQLSGV
						HVTEQSQTEGSWFGQHAYPF
25540	55908	Α	25681	568	724	RFSLGTVLDGIYPHQYDGYSR
1						WSAKRYL*WRNL*SGTHRSC*
						VNNFTRSNKK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25541	55909	ĪA	25682	1	974	MNGKLQSSDVKNETPYNIPLLI
23341	33,707	1	23002	1	,,,	NENVISSGISLISLWHTYADEHY
				1		RVIWPRDKKKPLIANSWVAVY
		1		<u> </u>		TVTGCGNSDRLNNIRTLRA\MA
		1				REFSIDVLDEMLQQFRVVTPDR
1				İ		PEHKQLPQHELAERQEKISTWL
						ELMKADGINPEELLGN\SSAAA
				1		
				1		PR\AGKKRQPGSAKYLFT\DVN
						GETKTWTGQGRTPKPIAQALAE
						DCHGKPEYGSAVHGCRCDDGS
						AGNLCFDLNRHCGLGVRLRPR
						KINGDYRSRFCHAGETGVCAG
						AETTQMAESGVSGAGLAVRGE
						RDGKRSPLQIAIMMNGSSGQAI
		1	2.500		1050	SSNVIILQLFLVEVIKSLRMLYM
25542	55910	A	25683	1	1359	
25543	55911	A	25684	72	918	PAPVAGAHCPLGLHRSSEHQEP
25544	55912	Α	25685	12	1090	`
						QLRHVARWPSS*SWLCHGTFR KLLPGKKRRGGNKKF*GRRAR
•						
						GVGCVVSRVWCGPACACSLRR
						WLAHTVHWVFTEAQSIKSHSS
						DMWPDGLALSPGCAMGPSGSC
1						SRGKNVGGATKNFRGVHNIYC
İ						HKPIPTTQHESKYARQTTTEEC
						TQTSSHDFRTTFSAVMQQRLQI
		-		Ī		ARNLVTHPKLVFMDEPTGGLD
						VSVQARLLDLLRGLVVELNLA
	1					VVIVTHDLGVARLLADRLLVM
						KQGQVVESGLTDRVLDDPHHP
<u> </u>						YTQLLVSSVLQN
25545	55913	Α	25686	1334	1545	RSQSRLERSCSVTGSM*RVSPA
1						RSFRRITKRVLAAARLAT*LPAS
	}					GRRTGVGVSVDSEFPSELLSSSF
		<u> </u>	2.50=	1.704	0101	TD
25546	55914	Α	25687	1784	2125	WLQHIHPQLLRCPVLRAQNVW
ļ						IRTLYDRH/RFVTRGTLGWIETG
						DF/DKVPPDLRFFAGGDRSI/RG
						YKYKSIAPKYANGLDIGKSPAA
						GAISPCLTFVMSARRGGHRGQS
	<u> </u>	 				ASGCRA
25547	55915	Α	25688	931	1213	NLHPAPAKRRAPARPQSRLAHR
						QRGRMRGAPRPFRQRQIKLCY
						ARCTPTICPTKVKSRSNTVTSG*
						TWSPRQRAKWWKSAKPPSAGL
						TSSARHP
25548	55916	Α	25689	999	1243	VDPVPWLPMALDITSTAPARQF
						RKRIIASLMPT/WGTSLRRYSVP
						PRRSGWKRWRFPPTGSTTCRPP
				<u></u>		FPAVCSSVCRLPATW

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25549	55917	A	25690	1225	2341	ARTFRLSNFGGSTRARGVYWD PNSDLCPTLDFSYKMPLLKNPL EQYYLVQGGFKRTDLNDTESD STTLVASRYWDLSSGWQRAIN LRWSLDHFTQGEITNTTMLFYP GVMISRTRSRGGLMPTWGDSQ RYSIDYSNTAWGSDVDFSVFQA QNVWIRTLYDRHRFVTRGTLG WIETGDFDKVPPDLRFFAGGDR SIRGYQY\KSIGPKLPNGD\LKG ASKLITGSLEYQYNVTGKCTK WQFGEWLAPGEVRLAPGATLA TPELVASCSTEGLNGLAANFHA ELRARLPWHGGAMKPRPVHLN TWEGFYFDLQPDKVKALATAA AALGVERFVLDDGWFNARHH
0.5.5.50	55010		25.601	45	222	DRAGLGDWWPDATKFPQGLG ELVAHVNPFYA
25550	55918	A	25691	45	233	SSYPRWWAAWWASTTARPSTR WRSSPR*SATTWASSPSPTSP*I HDRPGIGATHSSRFIPLK
25551	55919	В	25692	1	558	
25552	55920	A	25693	324	657	
25553	55921	A	25694	831	1244	WQPGGR*WSRQRSADGTGPDD LAAAYFPGDRFYRN*PSERC*H RRG
25554	55922	Α	25695	1	1215	
25555	55923	A	25696	1	840	
25556	55924	A	25697		1365	MPQNRGGFLDKKNPGAPLEGA VYVNCFTRFPGIGYEGRTAVFC GAAKVPPGLNYRGIGAFPGGNF ASKRSRGRNLAGARCLRFCPK\ WQ\NYSETASVQQQN*SVLARA IERGINAPLA\SSCGRFFDAVPA ALGCAPATLSYKGEAACALEA LAASCHGVTHPVTMPRVDNQL DLATFWQQWLNWQAPVNQRA WAFHDALAQGFAALMREQAT MRGITTLHPEGLERMSAGHES WLERRNGELGEGKGSKAYPTG DHKGADLYDDSNFLIQNIFVED TRGLLRQGREDLIERYKDPLNE NPKRCVEQLANWHKELEEYKK ASRIDIKPSREYASTIMNAIWTG EPSVIYGNVRNDGLIDNLPQGC CVEVACLVDANGIQPTKVGTLP SHLAAMMQTNINVQTLLTEAIL TENRDRVYHAAMMDPHTAAV LGIDEIYALVDDLIAAHGDWLP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25557	55925	A	25698	1	950	MPDEENRFNALKAVFSQLMTD ATLTPLFNYHYRISAPPGVNGL
	İ					SMKRAVVVFSGGQDSTTCLVQ
				1		ALQQYDEVHCVTFDYGQRHRA
						EIDVARELALKLGARAHKVLD
						VTLLNELAVSSLTRDSIPVPDYE
						PEADGIPNTFVPGRNILFLTLAA
						IYAYQVKAEAVITGVCETDFSG
						YPDCRDEFVKALNHAHKEKLD
						EDCKRRMYTNPLRVLDSKNPE
						VQALLNDAPALGDYLDEESRE
						HFAGLCKLLESAGIAYTVNQRL
]					AVNPEFKADPVVDIYLVASGA
						DTQSAAMALAERLRDELPGVK
						LMTNHGGGNFKKQFARADKW
						GARVAVVLGESEVANGTAVVK
						DLRSGEQTAVAQDSVAAHLRT
						KEKDSVEIYENENDQVEAVKRF
		1				FAENGKALAVGVILGVGALIG
		1				WRYWNSHQVDSARSASLAYQ
1						NAVTAVSEGKPDSIPAAEKFAA
						ENKNTYGALASLELAQQFVDK
						NELEKAAAQLQQGLADTSDEN
						LKAVINLRLARVQVQLKQADA
						ALKTL*IPD*RPDAGLRCKRSPR
				İ		SPAVCTDRQNAHEIPRPDSHLM PERR*EAPALPDFLNPARAAG
25558	55926	A	25699	1	3314	MKISRLGEAPDYRFSLANERTF
23338	33920	^	23033	•	3314	LAWIRTALGFLAAGVGLDQLA
		1				PDFATPVIRELLALLLCLFSGGL
						AMYGYLRWLRNEKAMRLKED
						LPYTNSLLIISLILMVVAVIVMG
						LRAMLMAMEPIWESDFHTLSY
						GFRPERSVHHAIRTVKLQLTDC
						GETRGRWVIEGDLSSYFDTVHH
						RLLMKAVRRRISDARFMTLLW
						KTIKAGHIDVEDKRAASEGVPQ
						NKIISPLLSNIMLNEFDQYLHER
						YLSGKARKDRWYWNN
25559	55927	Α	25700	997	1716	PYLTASSSAFQQLHLSGENHRD
						DGCAGYSRRTLAAQLQCHCEC
						DHNLPPDTPRLPRSGVVLPARF
		1				PRPLHTLISALHRRVLTRHARH
						ESSTLNPQRLPPAHRFFACQSRY
						VPGPAEYSYAVRSTDESG*A*W
						LRVPPNLIALTVQRSLPDLALPA
						QNCPPCRSMLEPAPYILLGMLA
		Ì				TWPACPG\HEVG*SHSAHPDEP
						ETLQTVSR*YRRFGRPVH*SVH
		_				VVGRYPHQACQCYLSSTAE
25560	55928	Α	25701	1	769	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25561	55929	A	25702	1	3707	MVRSAVVAGIAQAALPQSVVA
						WEYLVEDYDRIRNDIEAVLPEF
					ļ	ADYNQRIRHPGGFHLINAAAER
					}	RWMTPSGKANFITSKGLLEDPS
						SAFNSKLVMATVRSHDQYNTTI
				ŀ		YGMDDRYRGVFGQRDGYRDN
						CEEIAQLAEDCGNLFAVTVFRC
						PAGANLQGNRNVHCIDDLGKD
						RFNDIRNVQAAELLAEMAISSA
					ł	QRRMRAAKKVVRKKLTSGPAL
						PGKLADCTAQDLNRTELFLWK
						GEQNALLRGFPAVRLHSLL
25562	55930	A	25703	1	369	LPSHGAGLGTCSLPCLSLPPTP
						WAPVQPEPPRRAPPPAPWRPVP
						LTTQGLRN/DEHRVQDWQAAP
						PAAPLKVCKYTNQHPVFSTRFV
						NAPIDTLYLAALVRTWRTFMSS
						SRIVNTPIGTLYLAQ
25563	55931	A	25704	1	444	MRLHSSALGRSMGLGAVEQGA
				}		ALVGEAPAAQEPTEAGEGSGM
						AGCSPKACPAGRQLRPGPA\AA
						PSAGPPSPRPPGTPAGPQAPRAA
						PVPARASPSTPPCKLRERAAAF
İ						A/EPRKGLPQCSGGLKGSSSAA
		1				KVGAQAEEAPRASESCEDC
25564	55932	Α	25705	1767	2130	YTGQRLRRVYWVPQQCQPTCA
			}			ALDFSPSLSCLPA/AAGLRTYSQ
						PCLSLPPTPWAPVQPEPPQ*APP
		1				PAPRRP\AHRPPKG*GVQAHGA
					:	G\RQAAPPAAPVRDPLG\KPAGL
						LSLVGTWRTFMS
25565	55933	Α	25706	654	832	RRCARFLAGP*LPSHGAGLGTC
						SPPCLSLPPTPWAPVRPEPPRRA
		١.				PPPAPRRPVLCVFS
25566	55934	В	25707	1	537	
25567	55935	Α	25708	1	455	HCGSVIGGFLVSLTLRTKLRTL
		1				AVSVTALTVVCLEFIPSDVRMR
		1	1			SVSSFWWVRSLAGSRVKLQTF
						AVTVTALKAASLELFVPPGGLV
		l				VSLASGVKLQTAAAGK/RCLPV
		1				LRHAPALLSPWVVDGTGHRGA
1						GGGAHRGGSGRPGAHGGGGRL
25568	55936	Α	25709	1	314	SACREVWRERHEREPGLRGAL
		1				AGQLEFRVGVGLVRPALGGAG
						QPGAALDFSPGLSCLPA/AAGL
						GTCSPPCLSLPPPPWAPVQPEPP
						RRAPPPVHGAQSHRPPKG

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25569	55937	A	25710	144	519	HPGQRLRRVYWVPQQCQPIGA ELDFSPSLSCLPA/AAGLGTCSP PCLSLPPPPWAPVQPEPPQRAPP PAPW/WLSPIDHPRAEECECTA QDWQAALPAAPVWDPLEMSG LFDRAEKRYIRSFARKDSM
25570	55938	A	25711	271	743	ARTWNVPKVCVRIKANSHSIFT GLSCLPAGQGSGPAARHA*ASD PLRGLLCGPSLLDE/PPPPAPWR PVPSTTQGLRSAGEQRGTGRQL HLSAAPVRDPRGEAIWAPESGR SAASILKPARPPAHREERTTPDA PP*EL*HSPRRSAASLREDPWLH S
25571	55939	A	25712	1043	1348	GPARNRAQRRRAGTAGGPITPS AAAGPGAKPLIARRRQGWPAA PSAGPAKPTPTRISSWPAS\QPA AAVPARASPSTTLFKLREWAPA LASPERGSHSATVG
25572	55940	A	25713	22	226	NTRRWTEMTFDQVVRIFSIGNL QTVLQNRQPGGAIARCTGHIDP VTRFRPRAR*GSSHRNKAVDTQ RH
25573	55941	A_	25714	166	714	
25574	55942	A	25715	1	462	

	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25575	55943	A	25716	1	924	MFAIISYSSLAAVLLTATLTAAG
						IISFPVALCLVIGANLGSGLLAM
						LNNSAANAAARRVALGSLLFK
						LVGSLIILPFVHLLAETMGKLSL
						PKAELVIYFHVFYNLVRCLVML
						PFVDPMARFCKTIIRDEPELDTQ
į						LRPKHLDVSALDTPTLALANAA
						RETCALATPWTDDGRKYAYSA
						ASGGRRSATKVMVVVTDGESH
						DGSMLKAVIDQCNHDNILRFGI
						AVLGYLNRNALDTKNLIKEIKA
						IASIPTERYFFNVSDEAALLEKA
						GTLGEQIFSIEDMDLGDEVYTV
						GRPHPMIDPTLRNQLIADLGAK
						POVRVLLLDVVIGFGATADPAA
			:			SLVSAWQKACAARLDNQPLYA
					•	IATVTGTERDPQCRSQQIATLED
						AGIAVVSSLPEATLLAAALIHPL
						SPAAQQHTPSLLENVAVINIGLR
						SFALELQSASKPVVHYQWSPV
İ						AGQGKWLANPELLEADADAEY
				İ		AAVIDIDLADIKEPILCAPNDPD
						DARPLSAVQGEKIDEVFIGSCM
						TNIGHFRAAGKLLDAHKGQLPT
						RLWVAPPTRMDAAQLTEEGYY
						SVFGKSGARVSSIPCAVPCVWA
ŀ						RVADGATVVSTSTRNFPNRLGT
						GANVFLASAELAAVAALIGKLP
ļ.						TPEEYQTYVAQVDKTAVDTYR
						YLNFNQLSQYTEKADGLLKPRF
						RPWQRKILDTLATYHEQHRDEP
						GPGRERLRRMALPMEDEALVL
25576	55944	Α	25717	1	417	MGVLNNWLSEEESLWIQSRIHL
						RALRYYSNWRQYFAGYTFGRQ
						YWQSPEDDHLPLLREFLARYER
						PMKNCWKILDIEETTDVDIIRRA
						YLALLPSFHPETDPQDFKQLRQ
						AYEEALRIAQSPAKSVWQPEEY
						EVAEHEILLAFRALLASDSERFL
						PSAWQRFIQQLNYCSMEEIDEL
						RWSLCTIAMNTAHLSFECVVLL
						AERLRWLQEENTGEIDEEELGS
						FLYAIAKGNVFNFQTILHLPVA
						RRPDICTQHNDQSRCEGYQILS
						DKRGHQHGGGITALYQCGNAD
						PRAKGERLFFDAPA*CASSARV
						CVLSLAGASKKSRSPFARGSAL
						PHWYRAVMPPPCW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25577	55945	Α	25718	1680	2058	YRAYAPAFLPLSPSCPPLWRHC
						GSGRGENRRRDSSAHGPYAPRR
						QPDMPAHRQS*SFV/PASCSAC
ŀ		1				MDEPHKRLTVTAGTECGTSASS
		l				AALRAILKPCSSVCSSFFLLLLL
		1				PPPPPSSSSSSFFFLF
25578	55946	Α	25719	1115	3549	NPDRLLQLNLILETSK*QQMGD
						TRKKRS/NLTVAPIISPLKRYEQ
						HTELSRHINVVNLGDRNRRQQL
		İ				ILDQGKSSQRNIYLSHVAQKIPG
		l				RATYALSHSVDMITIQLGTLLR
						CAPPHRRRYAGRFFCRSAGNFL
ļ		İ				NVQGFDAVLVAVKHDLLLCLT
						IAPSLIVCTRLRSPWCGALRRCS
ŀ						TDGRSDLASSGVMFSIDTESAL
		1				TRWCLSLPHGALGKIIAEGLLS
						VIASKASAIVTTVAVVPVTRRSS
					}	LVNWHSGGSGLWRCNRTDERD
						QQASQGTTRMGRTQRNYSPTR
		į				QAIHKTGSPTMATTSNTEQPPH
						EKPNTLTQHPRNPDRTIHPKAT
		ŧ				HNELTQDTPQRDAKQNGEKTQ
}	,					RKKNCLSELNDAEDQAERFQD
						RLMLKLQVTTKPCSMTRLRDC
						AGIWSAANRWSGIGIDRMIMLF
		l				TNSHTIATLFSSRRCAHRNKFH
						LMNEAVRRLLRSSFRKSVLRRD
						ASFHDNQRVHSLQLLKERKKSL
						IHGRICAAKASLFQTRRYRWQK
		1				APLFSAQFRGRNRAVTHNRWV
						TTTTAIERTRARGLRPNSGRGF
						HSSPAWQLRHRKCRRCARRYH
		i				TVYWIKRRRSVARASTVVSGRI
						TSRRGENRRRDSSAIAHMLHAA
	,	1				KQTCLRIAVMIIRAASCSACMD
						EPPSDLTVTAGTDAGLAPAGYA
						LNERROLLHALTKSTALRTAHR
						GPNRTGWLNRRIVRFVMTALF
25579	55947	Α	25720	606	1562	
25580	55948	Α	25721	688	762	
25581	55949	Α	25722	492	737	SLPSGATLPPPALLLASAPDTA*
						WRCRER*YHLPAYAALSPAVA
						HRASLYSTLPSSVPWRRQCAGF
L						RARRSPEQAWAYPAR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25582	55950	A	25723	335	903	WVHDLPARKGSSARNIMLKSA PLRGHTRTRRKKMNIIEANVAT PDARVAITIARFNNFINDSLLEG
						AIDALKRIGQVKDENITVVWVP GAYELPLAAGALAKTGKYDAG IALGTGIRGGTAHFEYVAGGAS NGLAHVGQDSE\IPVAFGVLTT ESI/EQAIERAGTKVGNKGAEA
						ALTALEMINVLKAIKA
25583	55951	A	25724	3644	4545	SEAWPWRTAGTCLLCLNTT/AH VDKAIESRTVVADVLAKFENAL TGKLLTVSFQAHEAVIHGHHPA PEENLQVLAALRLQYLQGDYT LHAAIPPLEEVYSLQRLKARISQ STKTFTPCERLEKRRTSFLEGTL RRSFRTGSVVRQKVEEEQMLD MWIKEEVSSARASIIDKWRKFQ GMNQEQAMAKYMALIKEWPG YGSTLFDVECKEGGFPQELWL GVSADAVSVYKRGEGRPLEVF QYEHILSFGAPLANTYKIVVDE RELLFETSEVVDVAKLMKAYIS MIVKKRYSTTRSASSQGSSR
25584	55952	A	25725	1	735	
25585	55953	A	25726	147	504	STCPASAPPRPSAGPRILSGSSAS GSPLGSSSASSPAVHKG*RFWL TG*CGP*VSAGNK\LFERWDGS VQSIIPLQILESQGSKNLEVFFRG WMVAMDNRFVGLKGDSQQLS GQRIFKLG
25586	55954	Α	25727	1	639	
25587	55955	Α	25728	1	1247	

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25588	55956	Α	25729	1	2376	MAEVEEKKQRIFRKFTYRGVD
		İ				LGQLLDMSFEQLRQLYGARQL
						RRLNHGLWRNAEAQAAFAAE
	į					APMLGQEGGADHREAGSGEDA
						PAGYDHPSQDGGQQGGRLQRQ
						DLQPGGDQAGDDRPRPGRVLH
						HLQAQRGKKPM/ISTVTFGQ*P
						DPPLRAGQRL*RC/RLLIYGRGK
						CWALSGNP/RSGKTTLLKSISAR
·						LTPQQGEIHYENRSLYAMSEAD
						RRRLLRTEWGVVHQHPLDGLR
						RQVSAGGNIGERLMATGARHY
			1			GDIRATAQKWLEEVEIPANRID
ł				}		DLPTTFLRATHTVKMFSIPGYSS
						PGQLLDPETNINIGTSYLQYVY
İ						QQFGNNRIFSSAAYNAGPGRVA
ļ						TAIVLTLYIIYMVGFFATSGELT
						YEVWIGFFASAFTKVFTLLALF
						SILIHAWIGMWQVLTDYVKPLA
						LRLMLQLVIVVALVVYVIYGFV
						VVWGMRAALQISQSGQTCALL
						SKVFPTRSHTVSAQGGITVALG
1						NTHEDNWEWHMYDTVKGSDY
						IDAGRKRQEEGDWNSGVGDPS
						DAGDRKHQLKAAQLRPEAIKQ
						RRIANRLKRRTQFPRHLKGFVN
						TLKPIHGGCSVGTKCPQMASLP
						EGLHHPVKLVFLGKPGCLLTDR
						PLKGPNTAKPVAVDLNASATAI
						TSYSLYGGSMILLFLASTLYHAI
		ŀ				PHQRAKMWLKKFDHCAIYLLI
						AGTYTPFLLVGLDSPLARGLMI
25500	55057		0.5720	210	650	
25589	55957	Α	25730	219	652	SASWREAICLSRCRLPRSATAR
						GAGRGGCGLADRRASRQGKNL
						EAESAQEQNGHQHRIHESQHPC
						QGGAPVSHHQAAVRLRESQIQ
		1				GAAEKR*PTGDVIHPGQPVSGG
						/LK*YPSIPTCLLTTRTASLLKPP
2.55.5	4.50.50	ļ	2.550	1000	2100	GVRTSVCWQQLPR
25590	55958	Α	25731	1902	2100	RKFFSSSSGRS*PVRCSKL*SSID
						A*PLERIKRSRSYQVGSVGLCL
						RKSFHSTSAISAMPIGAPG
25591	55959	Α	25732	3223	3669	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		:	ļ	sequence		
25592	55960	Α	25733	1	4325	MPEPPTHSMGSCAAGASLTSTA
1					İ	PCSTAPSPIDHLRAKECERTAQ
		1				DWQAAPPAALVRDPLAALVRT
		l				WRTFMSSSRIVNTPIGTLCLAQ
						GFQWFSVLSQVHASTDQEIQE
						MHDEQANPQNAVGTLDVGLID
						SVCASDSPDRPNSFVIITANRVL
						HCNADTPEEMHHWITLLQRSK
						GDTRVEGQEFIVRGWLHKEVK
						NSPKMSSLKLKKRWFVLTHIPG
1						LLQEFREDALNWGPDEKIFKET
						ELVNDMDKINGRIERAE
25593	55961	A	25734	501	2871	TGCCCLRKTDCRRRESARGVH
23373	33701	 	25754	301	2071	AVYRRICARSMLLYRESVRPKA
						VTARVAVEAGIADYWYKYVG
			İ			LNGAIVGMTTFGESAPAELLFE
						EFGFTVDNPLASRAVRLRLPFN
						NDQVEAAVGWKLAVERHNGP
						TALILSRQNLAQVERTPDQVKE
			-			IARGGYVLKDSGGKPDIILIATG
		İ				
						SEMEITLQAAEKLAGEGRNVR
						VVSLPSTDIFDAQDEEYRESVLP
						SNVAARVAVEAGIADYWYKY
						VGLKGAIVGMTGYGESAPADK
1						LFPFFGFTAENIVAKAHKVLGV
i		1				KAGSHIPRKKYDVPGKKSFSPV
	İ					KYSTIGSPSPERPVSITTIPNSFVI
						ITANRVLHCNADTPEEMHHWIT
	ŀ	1				LLQRSKGDTRVEGQEFIVVEKL
		1				IRGLAMEDSRNMFALFEYNGH
						VDKAIESRTVVADVLAKFEKEP
]			VGKHLPTYDGQIAMGLESTAS
						VTRKHSSPSSLQNALSCETATSL
		1				IERRVRMALGTGDRRGLNTWL
			1			ARLPMEAKEKDEWRYWQADL
		1				LLERGREAEAKEILHQLMQQR
	1					GFYPMVAAQRIGEEYELKIDKA
		1				PQNVDSALTQGPEMARVRELM
		1				YWNLDNTARSEWAN\LLKNKS
		1				KPDRAHLPGYASTTQWWILSV
		1				QATI\AGNLGDHLKERFPLANN
		1				DLSKRYTSGKKIPQSYAMAIAR
		1				QKSAWNPKVKSPVGASGLMQI
1						MPGTATHTVKMFSIPGYSSPGQ
25594	55962	Α	25735	1	1195	
	1	<u> </u>	1	I		

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25595	55963	A	25736	1931	2676	FCGNTHQTYLGWVGGVANSEA *GEKWGTPDLVG/AERTETIQR LLGVCPEQGEGKIYIDGKQVDI RNCQQAIAQGIAMVPEDRKRD GIVPVMAVGKNITLAALNKFTG GISQLDDAAEQKCILESIQQLKV KTSSPDLAIGRLSGGNQQKAIL ARCLLLNPRILILDEPTRGIDIGA KYEIYKLINQLVQQGIAVIVISS ELPEVLGLSDRVLVMHEGKLK ANLINHNLTQEQVMEAALRSE HHVEKQSV
25596	55964	A	25737	1491	1667	TNGQWRAFYC*MRSGRNIRRR GEDISGGAGVSPRESPE*REVPE MGPGGAANRGTKRV
25597	55965	A	25738	419	823	VTLVAITRPIHVNRLRQYLVAIS LNETFGPSCKPSRSPCSSLSTCC WNAGVMHKK*VQGGRLCSYR N*SICLVVISPVVMGKRCIS*RC TAVLAALTVTVPSGVVAAHSV MLPRLPMKRSSIVPLPSNWRTS
25598	55966	A	25739	591	968	VKQCKQADEH*EQSLIPLERRR TTPLSVPQL/GKALTTVSFCASG GTSVNSLSHTS*ECMASAVAVK FTFPNLAVNFCTAAMAGPKVA LTSTLARIQAGSLARTGGRAYR WSRSTMLAKSFWCLMAL
25599	55967	A	25740	764	1005	QARPRAPRRAPYPILIRSSIITLIY /DGQRLGHLDDGTDSWNHVPT LNGQCTLVLIHEYVDTRGPEIK TILVISWWNILVFH
25600	55968	A	25741	463	786	
25601	55969	Α	25742	3	646	
25602	55970	A	25743		3008	MTMLQIVGALILLIAGFAILRLL FRALISTASALAGLILLCLFGPA LLAGYITERITRLFHIRWLAGVF LTIAGMIISFMWGLDGKHIALE AHTFDSVKFILTTALAGGLLAV PLQIKNIQQNGITPEDISKEINGY YCCFYTAFFLMACSDQRRLRG AMDKRYKFILVGFGEVFVFNLE SLDSGKKIHLLRRSHEEVMGSV MRTGTIDKGILPFIHHTRGNEIL FIPALFILFSLGGAVFGMGEEAV AFAIIIA

NO:	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
25603 55971 A 25744 2 3494				1	I .		
25603 55971 A 25744 2 3494 TITNLRPAAPGICEMAQGLSA GEYTGASIFVAEYSPDRRRGF GSWLDLRSIAEIELGAGGVOFF TIVGEANFLDWGWRIPFIFIAL GIIGLYLRHALEDTPAFQQHW KLEGODREGLODGPKVSLEK TKYWRSLLTCIGLVIATNVTY MLLTYMPSYLSHNLHYSEDH ALIIIAIMIGMLFVQPVMGILS RFGRRFVLLGSVALFVLAIP ILINSNVIGENLLPAMFPTHIR ALLLEMKN		sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
GEYTGASIFVAEYSPDRRRGI					sequence		
GEYTGASIFVAEYSPDRRRGI	25603	55971	ĪA	25744	2	3494	TITNLRPAAPGICEMAOGLSAG
GSWLDLRSIAEIELGAGGVGI TIVGEANFLDWGWRIPFTAG GIGLYLRHALEDTPAFQQHV KLEQGDREGLQDGPKVSLKE TKYWRSLLTCIGLVIATNVT MLLTYMPSYLSHNLHYSEDF ALIIIAIMIGMLFVQPVMGLLS REGREPFVLLGSVALFVLAPI LINSNVIGLNLLPAMFPTHIR ALLLEMKN 25604 55972	į						1
TIVGEANFLDWGWRIPFFIAL GIIGLYLRHALEDTPAFQUM KLEQGDREGLQDGPKVSLEK TKYWRSLLTCIGLVIATNVTY MLLTYMPSYLSINLHYSGEN ALIIIAMIGMILFVQPVMGLIS RFGRRPFVLLGSVALFVLAIP, ILINSNVIGLNLLPAMFPTHIR ALLLEMKN							
GIIGLYLRHALEDTPAFQQHV KLEQGOREGIQDGPKVSLET TKYWRSLLTCIGLVIATNVTY MLLTYMPSYLSHNIHYSEDF ALIIIAIMIGMLFVQPVMGLLS RFGRRPFVLLGSVALFVLAG RFGRRPFVLLGSVALFVLAG LILINSNVIGLNLLPAMFPTHIR ALLLEMKN 25604 55972							i l
KLEQGREGLQDGPKVS.KE TKYWRSLLTCIGLVIATNVT MLLTYMPSYLSHNLHYSEDF ALIIIAIMIGMLFVQPVMGLLS RFGRRPFVLLGSVALFVLAIP ILINSNVIGLNLLPAMFPTHIR ALLLEMKN							i I
TKYWRSLLTCIGLVIATNVTY MLLTYMPSYLSHIALHYSEDH ALIIIAIMIGMIGHVQPVMGLLS RFGRRPFVLLGSVALFVLAIP, ILINSNVIGLNLLPAMFPTHIR ALLLEMKN 25604 55972		ŀ					``
MILLTYMPSYLSHNLHYSEDH ALIIIAIMIGMLFVQPVMGLLS RFGRRPFVLLGVALPVALIP. ILINSNVIGLNLPAMFPTHIR ALLLEMKN 25604 55972							t i
ALIIIAIMIGMLFVQPVMGLLS RFGRRPPVLLGSVALFVLAIP, ILINSNVIGLNLLPAMFPTHIR ALLLEMKN 25604 55972							I I
RFGRRPFVLLGSVALFVLAIP. ILINSNVIGLNLPAMFPTHIR ALLLEMKN							1
ILINSNVIGLNLLPAMFPTHIR ALLLEMKN							'
ALLLEMKN			i				1
25604 55972		ı					
25606 55974	25604	55972	Α	25745	1	277	
25607 55975	25605	55973	A	25746	902	1006	
25608 55976	25606	55974	A	25747	2	1265	
GHLRPVERQCSPDKRRFIAAI	25607	55975	Α	25748	409	525	
REIRGKQRAGHIFQLLSRCLL	25608	55976	A	25749	91	431	NHKPGNIDVARRIQRGFAGDQI
LNHCQRRAAHFRFQLSNQRH QLLPV/HYHAAEREYPAGACI RWLL			İ				GHLRPVERQCSPDKRRFIAADG
QLLPV/HYHAAEREYPAGACIRWLL	i						REIRGKQRAGHIFQLLSRCLLQI
RWLL							LNHCQRRAAHFRFQLSNQRHQ
25609 55977							QLLPV/HYHAAEREYPAGACLV
25610 55978		:					RWLL
25611 55979 A 25752 I 2786 MAAVIEQIRRAVLALVTGVV GDIPVVWDIVWNATAAFIAV SLLLDESGFFEWAALHVLRW YGPGRLPFNRIGLIRRITRRIRG LLFGLVTFILLLNRSQNPAQD LLAFRARTGKQTAQFIHHLPR VFTDKTGSSNGLAPQQCSAGELILNEKLAKQLVTAANWV MQSDEGEINPVDILRWPGVM AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPF LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H./RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAVGYFKVHVLG*RSVTPVTDIVKLEFTRLRLPGYTKSIE						I	
GDIPVVWDIVWNATAAFIAV SLLLDESGFFEWAALHVLRW YGPGRLPFNRIGLIRRITRRIRG LLFGLVTFILLLNRSQNPAQD LLAFRARTGKQTAQFIHHLPR VFTDKTGSSNGLAPQQCSAG GELILNEKLAKQLVTAANWV MQSDEGEINPVDILRWPGVM AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPE LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H /RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAVG YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE			 				
SLLLDESGFFEWAALHVLRW YGPGRLPFNRIGLIRRITRRIRG LLFGLVTFILLLNRSQNPAQD LLAFRARTGKQTAQFIHHLPR VFTDKTGSSNGLAPQGQCSAGELINEKLAKQLVTAANWV MQSDEGEINPVDILRWPGVM AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPE LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H /RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAVG YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE	25611	55979	A	25752	1	2786	1
YGPGRLPFNRIGLIRRITRRIRG LLFGLVTFILLLNRSQNPAQD LLAFRARTGKQTAQFIHHLPR VFTDKTGSSNGLAPQGQCSAG GELILNEKLAKQLVTAANWV MQSDEGEINPVDILRWPGVM AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPE LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H //RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE]						
LLFGLVTFILLLNRSQNPAQD LLAFRARTGKQTAQFIHHLPR VFTDKTGSSNGLAPQGQCSAG GELILNEKLAKQLVTAANWV MQSDEGEINPVDILRWPGVM AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSF APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPF LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H /RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAVG YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE	1		1		1		
LLAFRARTGKQTAQFIHHLPR VFTDKTGSSNGLAPQGQCSAGELILNEKLAKQLVTAANWV MQSDEGEINPVDILRWPGVM AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPE LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H //RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV/G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE	1		1		1		· · · · · · · · · · · · · · · · · · ·
VFTDKTGSSNGLAPQGQCSAGELILNEKLAKQLVTAANWVMQSDEGEINPVDILRWPGVMAQEQDLDAIAAEILAALDGTIDFIVARETEGQSGLKRVYHSFAPDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRWVSSSPLETWNKRWAKISPAPFLLPISRLEKSEGDWLPETVISANMPSSLSLIISSL*FKVRDV*HVRI*TLRGHYRVINWPNFNIVVQGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\GVFKVHVLG*RSVTPVTDIVKLEFTRLRLPGYTKSIE							1
GELILNEKLAKQLVTAANWV MQSDEGEINPVDILRWPGVM AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPE LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H //RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							
MQSDEGEINPVDILRWPGVM AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPF LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H /RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							
AQEQDLDAIAAEILAALDGTI DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW VSSSPLETWNKRWAKISPAPE LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H //RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							·
DFIVARETEGQSGLKRVYHSE APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW. VSSSPLETWNKRWAKISPAPE LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H. /RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\GYFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							1 -
APDIREFTRDAIP 25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW. VSSSPLETWNKRWAKISPAPF LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H //RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							
25612 55980 A 25753 5 402 EIFSVVWIIMTRGDVVTISRW. VSSSPLETWNKRWAKISPAPF LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H. /RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							DFIVARETEGQSGLKRVYHSPG
VSSSPLETWNKRWAKISPAPF LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H /RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							
LLPISRLEKSEGDWLPETVISA NMPSSLSLIISSL*FKVRDV*H /RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE	25612	55980	Α	25753	5	402	EIFSVVWIIMTRGDVVTISRWR
NMPSSLSLIISSL*FKVRDV*H //RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE					1		VSSSPLETWNKRWAKISPAPFS
/RI*TLRGHYRVINWPNFNIVV QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							LLPISRLEKSEGDWLPETVISAF
QGIGKRLANSWLVFTLFHRI 25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE					İ		NMPSSLSLIISSL*FKVRDV*HFF
25613 55981 A 25754 48 219 PETCGHLWAYVWPSCAAV\G YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							/RI*TLRGHYRVINWPNFNIVVS
YFKVHVLG*RSVTPVTDIVKL EFTRLRLPGYTKSIE							
EFTRLRLPGYTKSIE	25613	55981	Α	25754	48	219	PETCGHLWAYVWPSCAAV\GL
							YFKVHVLG*RSVTPVTDIVKLL
	L						EFTRLRLPGYTKSIE
25614 55982 A 25755 1 912	25614	55982	Α	25755	1	912	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	<u>.</u>	
25615	55983	A	25756	1	1857	MRVCARACVRTRTRMCVYAH
23013	33703	ľ` .	23730	•	1037	TCVCVRTYAYVRVRVRVHVRA
		1				RVRVCAYARTRVRNSLSILPFIQ
		1				LTLATPIHHIHQEEFNIRGIVPVL
	,	1				RRVKPDLAIGIDITPSCDTPDLH
						DYSEVRINQGVGITCLNYHGRG
						TLAGLITPPRLIRMLEQTALEHN
						IPVQREVAPGVITETGYIQLFLP
						GWEIGFSPLALLLAFLCSTSPGF
						GDPDGLGVIAYQDTVRPNAAT
	'	İ				
						AISELNALAVKGVILTGDNPRA AAAIAGELGLEFKAGLLPEDKV
						KAVTELNQHAPLAMVGDGIND
						APAMKAAAIGIAMGSGTDVAL
		İ				ETADAALTHNHLRGLVQMIEL
						ARATHANIRQNITIALGLKGIFL VTTLDDRVVAGSAGRYGGDGA
						i
		İ				GDSECVKIVAQEIRQTDRNIER
						DRSPRPSGERVRVRGKGGIEAN
						QPLSTAFTNQITIRQSIRLFSNQF
						VFHKGISRVVAGSSAIISNVSPG
		1				CILRTPNCRRGFPISCTGEGKAA
		1				YGWQRDGGHAEYLLAEEKDLI
						LLPDALSYEDGAFISCGVGTAY
						EGILRGEVSGSDNVLVVGLGPV GMMAMMLAKGRGAKRIIGVD
						MLPERLAMAKQLGVMDHGY*I
			ŀ			LPDVY/V*IGVARVSWMNGRID
						SELRTRVRAYAHTRTRARTCTR
	İ					TRTRTYAYVRTHTHVCAYTHM
						RVRVRTHARAHTR
25616	55984	A	25757	3	1180	SKLGTRRSVVWA*SPSTSPTLW
23010	33904	^	23737]	1100	CSTFSAAGHSSMKRMNEFVDL
						LPAQQRMKGENWYRGTADAV
						TQNLDIIRRYKAEYVVILAGDHI
						YKQDYSRMLIDHVEKGARCTV
						ACMPVPIEEASAFGVMAVDEN
						DKIIEFVEKPANPPSMPNDPSKS
						LASMGIYVFDADYLYELLEEDD
	-					RDENSSHDFGKDLIPKITEAGLA
						YAHPFPLSCVQSDPDAEPYLAR
						CGYAGNFMESEPRSGLIRVSCM
						PAGPPLPLPAARSEAPKAAGTV
				1		ASVPSIAPARLRPVPDPVELVLV
				1		AAEFITPGDPTPRLHGSGFIDIR
						QIHHQTRSHLEGVKTGIRFLNH
						FSGNPQGGIAHVNGVARFQVK
						QCHQAWGQQYAARLRFQARGI
25617	55985	A	25758	16	543	SLQIAIHRVDIIHRFDVRQL
25618	55986	A	25759	1	3387	
23010	122300	<u> </u>	23133	<u>l' </u>	3301	<u></u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25619	55987	Α	25760	1688	2234	LSVGKTFPVLVPAAFRKVSATG
						AAVPGTPSILALPSYSTTPFGVC
						TSASMNVPVLTLRIRTRVTGEV
		1				AAIKPFSMANAPTPDSILPQLGV
		ŀ				VSTRCS*TITCANR*STSARGSL
						ERLIMATLL/WVAGTFD*SVWF
						GHDTDVVTLYRFHEALCYSVT
						LRTPHRRVLRFKSQHPGELACFI SPVA
25620	55988	Α	25761	905	4762	PPHNWMPSNATPGIAFVWCAY
						GAI/LPGDAPVPVVDDYRKVVR
		Ì				KDTKGLIARWKYFWMSVIALG
		ļ	Ī			VAFALYLAGKDTPATQLVVPFF
						KDVMPQLGLFYILLAYFVIVGT
	į					GNAVNLTDGLDGLAIMPTVFV
						AGGFALVAWATGNMNFASYL
						HIPYLRHAGELVIVCTAIVGAG
						LGFLWFNTYPAQVFMGDVGSL
						ALGGALGIIAVLLRQEFLLVIM
						GGVFVVETLSVILQVGSFKLRG
		<u> </u>				QRIFRMAPIHHHYELKGWP
25621	55989	Α	25762]1	1448	MCHVLLAGRQRRTAGPEHALT
						PPECGSAAIGPAVYKWAHATG
						VMIPDRPRSPHAGAEERHWHH
						PQPELISQRLGSFSSGPTKYIHEF
						QYLTLSYNLTWSDLNVILTFILP
		İ				LDEWERVFSLAQSHTDNCRLH
]				KPDLQEGIREVPQEDPQWNYQ
		İ				ENSPGIARRDYMISCLVEGLKK
		1				AAYKAINYDKLKETTQGTDKN
			ł			PAQSMACLAATMRHFAALDPE
		l				ALAPHRDVNHGIKLTCKTIRSSI
						DISREIRAPRECRSTEMASWHPP
						ADLTECPVESGYSVLSGRTDCQ YSHTVRRACLTTIDSRPAVRQL
	İ	l				LKIREVQLLSLLRGCVRSRRSGS
						~
						INAQYCRCLERVVTLPFFDSGPS LMNHAPHLYFAWQQLVEKSQL
						MLRLATEEQWDELIASEMAYV
						,
						NAVQEIAHLTEEVDPSTTMQEQ
						LRPMLRLILDNESKVKQVLQIR
						Q\DGLAKLVEQLRHLSKKHHLD
	J		<u></u>		L	TDSTSSSLPGVRTPYSPIRGGEE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25622	55990	A	25763	1029	1665	RPESISSVASKTRLGRSFHGAAR RTREYHSSTCNASIAAAATV\A AQGHPDTTSTPPASVDTTATNT QTPAVTAPAPAVDPQQNAVVS PSQANVDTAATPAPTAATTPDG AAPLPTDQAGVTTPVADPNAL VMNFTADCWLEVTDATGKKLF SGMQRKDGNLNLTGQAPYKLK IGAPAAVQIQYQGKPVDLSRFIR TNQVARLTLNAEQSPAQ
25623	55991	Α	25764	658	1307	SNRLYTDVESINEIEGIYMKLRF ISSALAAALFAATGSYAAVVDG GTIHFEGELVNAACSVNTDSAD QVVTLGQYRTDIFNAVGNTSAL IPFTIQLNDCDPVVAANAAVAF SGQADAINDNLLAIASSTNTTT ATGVGIEILDNTSAILKPDGNSF STN\QNLIPGT\NVLHGLAFRAR CVSTANLRVPRPICFDPGSSYQA NTIGSGRCADCQPDA
25624	55992	Α	25765	167	232	
25625	55993	A	25766	441	686	
25626	55994	A	25767	206	331	
25627	55995	A	25768	674	900	VVPICTMLEEALERQNLTASLC G*LKKACGRSWQTSPIKRSLNG AKFSRTSIIMGFITAEKWSGWEI WSARNVTS
25628	55996	Α	25769	41	222	
25629	55997	A	25770	17	525	ESLECSGPSWNASGTQSGFYCH RIAPILPEKRLARMVEKTWGLG LRRRTGFRRQNLGGGTGTLVE CTSENS\KPSKSSFRKQRRGQRR EGH*SLRR*IRRERTERMQASR VTWNSFPQVPAAKKPSSLSESG *P*ATNYTREQDSMWNCVHHY WQPPTSVSRLVHLNSG
25630 25631	55998	A	25771 25772	698	2204	PYLRIRLTALISRSSTVMRILFAI SKRPIT*PKRPKPTTMTWGLSSL THGFSSSSSGPLNWRDSHLSTSF ISNGVVIMESVTVTSNNAIWSPS SSVAAVNTSRTPEAAIKRNTGR SK

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25632	56000	A	25773	3	891	RYRNQDDHELQIHTGTRFYVEL
						CGFRVNPPWLATDVTHAAMAL
		l				LTLVTTGLHIVGEWGKHATPPV
						PYSICSNPPRRAAYITVWRNRPC
		1				TLGDHLFSIAYPRGLRYDMGG
						GGEGTAWANVRQALARTTGR
İ		l		•		AQHWTKDNAKSSREARHPRQN
		i				KTKTHLGKPVLRWV/DPSFDQS
		l				KYDSIVWNPI/TYYPVPKPSTQV
						GQKV/LDKILNYTNTEMKEAIA
						H/DTSKEGLQFYEVVPVALVVA
						GTQMATGLRTMDTRLYFEGELI
						DAAPNKPVIKVVRQGEEPDWP
		ľ				SKGDAGDGEKVPSSLSESG
25633	56001	Α	25774	1	285	
25634	56002	Α	25775	1	302	SFSSFTQRTASVIISSTSTQSRLA
						TSPILAIARARTSRRCPEKPQLG
						YPDSAAHVFQQDRATKYVINR
						NSKKP*ICSACRSTVSTRQHRR
						WRGSLQPLSQ
25635	56003	Α	25776	307	473	
25636	56004	В	25777	67	809	
25637	56005	A	25778	44	301	
25638	56006	A	25779	1	1519	MINRLQHTVYTHKWNSYRTEP
		ŀ				DRRHTALRNSFCLVRLNDSSHS
		ł				RSLDSKQGHTEITFAATEHPPSP
		1				VSLRIHDVAFDDGRQTSRAPCF NLEQPGKLDCTKRLISDCWVM
						HPGESWHGFKDIPDNWSMLDPI
		ŀ				KISILAPGMGEDGELEETGVPA
						ALVTAWLGRHGIVPTRTTDFQI
						MFLFSMGVTRGKWGTLVNTLC
						SFKRHYDANTPLAQVMPELVE
						QYPDTYANMGIHDLGDTMFA
					}	WLKENYSGARLNEAYS/GLPSII
						ERHVVDAQGYRGGGMLCSCES
						NFCVSLLRIQAS
25639	56007	A	25780	461	789	HHVLSELQQLYTAYRSVDHIID
		-				HSLAIARAPQRRRCPEKPQLGY
						PDSLQHVFQQDRATKYVINGTS
						KKP*ICSACRSTVTHGHTDAGE
						EVWNPFAVLVPSGKDARVWG
25640	56008	В	25781	1	990	
25641	56009	A	25782	1	3066	
25642	56010	Α	25783	583	904	WGIRTTDFQIMFLFSMGVTRGK
						WGTLVNTLCSFKRHYDANTPL
						AQVMPELVEQYPGR*FTGVRK
						KSNN*FRQFGGVDTAGNNL\SEI
						FSASASAEIFPANSFCNRLA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or peptide sequence	deterion,possible nucleotide insertion)
25643	56011	Α	25784	331	660	TGIQLPGAP*SQPWSTRRRAVV
						RPLYRVN*AVP*NR*VIRSKRA
						MPKRTDIKSILILGAGPIVIGQRL
						GWRDSDGVRLCGVLGSQENVC
	-					ELDEKGEERWSISGTSSEFTR
25644	56012	A	25785	1	798	MSDAEDGARTTLGSGNAVERL
		1				ADALSQQNVTVIKSTSLDDGFA
						ILSSNEAIDCLMFSYQMEHPDE
		İ				HQNVRQLIGKLHERQLNVPVFL
						LGDREKALAAMDRDLLELVD\
						EFAWI\LEDTADFIAG\RAVAAM
:						TRYRQQLLPPLFSALMKYSDI/Q
						SEYSWAAPGHQGGVGF\TKTPA
						GRFYHDYYRLLRVAWVLHKTE
		l				CRLINFPETPIFGNSFRYDIEVSN
ļ		ŀ				ESPDEEVKLRRHHLARCMKNF
					}	KTDIYFVSTFEPSTKSVDLLTVE
						TFAGT
25645	56013	A	25786	211	414	RVCECIQLRRIEIFRFMISSQPNL
						PEDLHPQIIAKPSQKRLIEQ*SSK
						LTVTKARRQGSRYKSSRFFS
25646	56014	Α	25787	1694	4974	LSLLSAK/PVYMVPSRNRYGIIG
		ŀ				PI/YPQEMQPETLQKKISE/SPLT
						KDKAGQKPSYCVV/TNCTYDG
						VCYNAKEAQD/LLEKTSDRLHF
						DEAWYG/YARFNPIYADHYAM
						RG/EPGDHNGPTASYIHVREGV
						GRLNFFRFNQAYMMH/ATTSPL
						YAICASNDVAV/SMMDGNSGLS
				-		LTQEDCWVMHPGESW/HGFKD
		l	[IPDNWSMLDPIK/VSILAPGMGE
						DGELEE/TGVPAALYSNLCLSSG
						TTGLSTEALLE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25647	56015	A	25788	11	1709	MSKPKYPFEKRLEVVNHYFTT
25011				ľ		DDGYRIISARFGVPRTQVRTWV
						ALYEKHGEKGLIPKPKGVSADP
				1		ELRIKVVKAVIEQHMSLNQAA
						AHFMLAGSGSVARWLKVYEER
						GEAGLRALKIGTKRNIAISVDPE
						KAASALELSKDRRIEDLERQVR
						FLETRLMYLKKLKALAHPTKK
1	l					LSASLTSMRTLKSVFSEIYHENR
						GRYGYRRVTLSLHREGKQINH
		1				KAVQRLMGTLSLKAAIKVKRY
						RSYRGEVGQTAPNVLQRDFKA
						TRPNEKWYTDVTEFAVNGRKL
		1				YLSPVIDLFNNEVISYSLSERPV
		1				MNMVENMLDQAFKKLNPHEH
						PVLHSDQGWQYRMRRYQNILK
		1				EHGIKQSMSRKGNCLDNAVVE
		ł				CFFGTLKSECFYLDEFSNISELK
		[DAVTEYIEYYNSRRISLKLKGL
			•			TPIEYRNQTYMPRVNCPLFDTL
		ŀ				RIPDIGGVVRARAIAKLLNDTD
						MAIIDKRRPRANVSQVMHIIGD
	1					VAGRDCVLVDDMIDTGGTLCK
						AAEALKERGAKRVFAYATHPIF
						SGNAANNLRNSVIDEVV/DLRY
						HSAER*NQITAERAYSDPVRKI
25648	56016	A	25789	2902	3271	SSAARHCAGCGDVRSVCRSGR
						ADAPRMHPGESWHGFKDIPDN
						WSMLDPIKISILAPGMGEDGE/P
						TRTTDFQIMFLFSMGVTRGKW
						GTLVNTLCSFKRHYDANTPLA
						QVMPELVEQYPAFVINQ
25649	56017	Α	25790	185	663	SPVPSSRPSTRHRLPDGSSSSQFS
						SAIAARKVVNSCILCSKICQN*F
		İ				NRRAASPLKRLP*SGASGMIT\P
						NQPAPAQ/PAK*LYTRLDEVKP*
						*KITHGNGVSFCPTG*A*RTGIS
						ASSRDLEVQSKKRSFPALNWPA
						VYCCSALGSAPKCKPIPFLRAT
		L_				V

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25650	56018	Α	25791	684	4208	NGLFWLFQPWLFADETRRRLE
25050	30010	1			1200	ERFRDRYQIIHHSSQVCHSDQA
		1				VYVLTQERVNERDDIVDIDLFV
		1				IDEFYKLAFRQLKSGDIDHQDE
						RVIELNIALSKLLKVSRQFYLTG
						PFVNSIRGLEKLGYPHTFVSTDF
						NTVALDVKTSASKRMTTKPSS
						KRWGKSRTCVDADHIKNGIGL
]						HFGALPRALQQYTADQFNAGK
						LRFLLCTSTHEGVNTIAKNVVIY
						DNRAGSTRTISADIINSVICGLN
						TLKSFDTRWALL
25651	56019	Α	25792	1	896	
25652	56020	Α	25793	231	721	IITAEKGRSREQQLAIL\PLNLAD
						MRQLQQQHSQTRLADTAADGL
1						RHFTAQQRLMPLQLQAIFVTGQ
1						RQLMLQRVGY/R/LHGSPWRKA
ŀ		İ				QTHLPVPGSTPEYRRSDPHGPA
		ł				LTGCSSHHNPVRARRGMTIAQL
						AANPDKEYRARSFAIARSRPLA
						SRSGYETVTPGC
25653	56021	Α	25794	1	933	MLEAAVCLCVFVSCSLAEGRA
						CCRVAVSIPVSVRARIHLLESDH
						LGRFREPKSIPRKDAQMDTFRT
						MVREPATFLDQVDGLQQSDDIL
						RLLPPELATLGITELEYEFYRRL
						VEKQLLTYRLHGESWREKVIER
		1				PVVHKDYDEQPRGPFIVCVDTS
						GSMGGFNEQCAKAFCLALMLI
						AIANKWRCY\IMLSS\TEIVRYE
						LSGPQGIEQAIRFLSQQFRGGTD
						LASCFRAIMERLQSREWFDADA
						VVISDFIAQRLPDDVTSKVKEL
						QRVHQHRFHAVAMSAHGKPGI
0.5.5.1	56022		05705	102	1100	MRIFDHIWRFDTGMRSRLLRR
25654	56022	C	25795	83	1108	
25655	56023	Α	25796	1	747	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25656	56024	Α	25797	25	1741	IMERLKRMSVFAKVVEFGSFTA
						AARQLQMSVSSISQTVSKLEDE
1						LQVKLLNRSTRSIGLTEAGRIYY
						QGCRRMLHEVQDV\HEHLYAL
						NNTPIGTLRIGCSSTMAQNVLA
						GLTAKMLKEYPGLSVNLVTGIP
						APDLIADGLDVVIRVGALQDSS
						LFSRRLGAMPMVVCAAKSYLT
						QYGIPEKPADLSSHSWLEYSVR
						TDNEFEL\IAPEGISTRLIPQGRF
						VTNDPMTLVRWLTAGAGIAYV
						PLMWVINEINRGELEILLPRYQS
						DPRPVYALYTEKDKLPLKVQV
						VINSLTDYFVETPTPLSSLTSLPT
						IEICCMVSIPGTNQRLRPFTGFA
						THRLQSDLLLQVMTISRGRFSR
						SARFNGPTPSGRRLIRVIISTESG
		1				SRSSNPRTTGCFEEHCDHYRAII
		ł				DSADGLPMVVYNIPALSGVKLT
						LDQINTLVTLPGVGALKQTSGD
						LYQMEQIRREHPDLVLYNGYD
						EIFASGLLAGADGGIGSTYNIM
						GWRYQGIVKALKEGDIQTAQK
	1					LQTECNKVIDLLIKTGVFRGLK
						TVLHYMDVVSVPLCRKPFGPV
						DEKYLPELKALAQQLMQERG
25657	56025	Α	25798	1	1005	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	}	
25658	56026	Α	25799	1	2392	MEMVAASKMRKSQDRMAASR
						PYAETMRKVIGHLAHGNLEYK
						HPYLEDRDVKRVGYLVVSTDR
						GLCGGLNINLFKKLLAEMKTW
						TDKGVQCDLAMIGSKGVSFFNS
						VGGNVVAQVTGMGDNPSLSEL
						IGPVKVMLQAYDEGRLDKLYI
						VSNKFINTMSQVPTISQLLPLPA
						SDDDDLKHKSWDYLYEPDPKA
						LLEQCSCMLRSPAWGITSTDTT
						RARTDLACPIWLLLDPYPKLMG
		ŀ				NAPKALASVDKRGVPVNTILVS
						ALVTALCVLINYLAPESAFGLL
						MALVVSALVINWAMISLAHMK
						FRRAKQEQGVVTRFPALLYPLG
						NWICLLFMAAVLVIMLMTPGM
						AISVYLIPPLQTNTFNSTTIAMA
						DYQMESLSAEINFAAAKLARA
						CADEWTARTPEKPRYVAGVLG
						PTNRTASISPDVNDPAFRNITFD
						GLVAAYRESTKALVEGGADLIL
		1				IETVFDTLNAKAAVFAVKTEFE
						ALGVELPIMISGTITDASGRTLS
		1				GQTTEAFYNSLRHAEALTFGLN
		1				CALGPDELRQYVQELSRIAECY
5						VTAHPNAGLPNAFGEYDLDAD
						TMAKQIREWAQAGFLNIVRGC
						CGTTPQHIAEMSPGRKRLTRAE
		1				SSSVPSTLPGKAWLEHALPLIAE
						QL/QGRRSAVFIPF\AGVTQTWD
						DYTAKTAAVLAPLGVSVTGIHS
						VVDPVAAIENAEIVIVGGGNTF
25659	56027	Α	25800	412	1115	LAGEVMDLLLLSNSTLPGKAW
						LEHALPLIAEQLQGRRSAVFIPF
						AGVTQTWDDYTAKTVAVLAPL
						GISVTGIHSVVDPVAAIENAEIVI
						VGGGNTFQLLKQCRERGLLAPI
						TDVVKRGALYIGWSAGANLAC
						PTIRTTNDMPIVDPQGFDALNLF
						PLQINPHFTNALPEGHKGDPRL
						FR\IRELLVVAPELTIIGLPEGNW
						ITVSKGHATLGGPNTTYVFKAG
		1		<u> </u>		EEAVPLEAGHRF
25660	56028	Α	25801	1	360	MDHAIAGIAVQFRGGNLLRFK
						AFQQRVKYRMCQAERAINRLF
						NIAVKRLTGNRLDDKSKQHIIDI
						AVDIPKWSDHLHIIRRKHHDPT
1						QDSGTHLPAVADDGFRTSVRN
						RSA*HIRYLTRC*KALNRNKLPP
}						RN*TAIPAMA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	• · · · · · · · · · · · · · · · · · · ·
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
25661	56029	A	25802	1032	1233	LVQFLRICFHILQRF/LPSH*RQK
						DESLRALPVSFLCWRTDSDFPA
		ļ				DISYQNYR*YNREPYLPRPDSG
25662	56030	Α	25803	1	1394	MTRDGLANKALAVARTLADSP
						EIRQGLQKKPQESGIQAIAEAVR
						KRNDLLFIVVTDMQSLRYSHPE
						AQRIGQPFKGDDILKALNGEEN
						VAINRGFLAQALRVFTPIYDEN
						HKQIGVINDSRWSIIWSVLFGM
						LVGLIGTCILVKVLKKILFGLEP
						YEISTLFEQRQAMLQSIKEGVV
	İ					AVDDRGEVTLINDAAQELLNY
						RKSQDDEKLSTLSHSWSQVVD
	İ					VSEVLRDGTPRRDEEITIKDRLL
						LINTVPVRSNGVIIGAISTFRDKT
						EVRKLMQRLDGLVNYADALRE
						RSRLEHILHNQRVPFIMKLHHR
						MLRHFIAASVIVLTSSFLIFELV
						ASDRAMSAYLRYIVQKADSSFL
		l				YDKYQNQSIAAHVMRALAAEQ
					:	SEVSPEQRRAICEAFESANNTH
						GLNLTAHK\SRAYAAHYKPHPL
		ļ				TATQLWK\RSTITRF*SGSGRQP
						PPG*LRFRSWDGRREISLLSRSL
25663	56031	A	25804	3	219	
25664	56032	Α	25805	14	265	
25665	56033	A	25806	744	1566	FASWLRVFSITNCAAPVTSESC
						ALDLGSAEAKAWIGVENPHRA
						DVLTELRRSTVARVCTGRAGPR
						PRTQALLRFLADHSRSKDTVLK
						EVPEEWVKAQGLLEVRSEISDK
						NLYLTRPDMGRRLCAEAVEAL
		ŀ				KAQCVANPDVQVVISDGLSTD
		İ				AITVNYEEILPPLMAGLKQAGL
						KVGTPFFVRYGRVKIEDQIGEIL
}						GAKVVILLVGERPGLGQSESLS
	1	l				CYAVYSPRMATTVEADRTCIY
]				NIHQGGTPPVEAAAVIVDLAKR
						MLEQKAF\GINMTR
25666	56034	A	25807	596	745	
25667	56035	Α	25808	151	660	STTERLGDDFVRAANIILHCEG
		1				KVVVSGIGKSGHIGKKIAATLA
ŀ		1				STGTPAFFVHPAEALHGDLGMI
		1				ESRDVMLFISYSGGAKELDLIIP
						RLEDKSIALAPNSRTVIT*CWFS
						AGDGGLPGPDYGQDFAAHPAG
						AGALYKCRDLINDNILRTVSCQ
						SHDDGDIRGMAFARQ
25668	56036	Α	25809	314	583	
25669	56037	Α	25810	i	2982	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
<u> </u>				sequence		
25670	56038	Α	25811	120	790	EGLKTKGEKASPPSCLSFCKNE
		1				FVISSTDYALSQSQSGGVMHRQ
		l				SFFLVPLICLSSALWAAPATVN
		l				VEVLQDKLDHPWALAFLPDNH
1		1				GMLITLRGGELRHWQAGKGLS
						APLSGVPDVWAHGQGGLLDVV
						*APDFAQSRRIWLSYSEVGDDG
		l				KAGTAVGYGRLSDDLSKVTDF
						RAVFRQMPKLSTGNHFGGRLV
						FDGKGYLFIALGENNQRPTAQD
						HRVPSWYK
25671	56039	Α	25812	707	1141	ASHTATSPRPVRLSSSIASITLAV
						NATCGMASSRRIR*CTLFSKRA
						PSAPAGWERAKSSSLNPRACMT
						AIASASPIIRVLTVLEVGAKCIG
						QGSRSDSTELSPFIKMVPPNSLT
ļ		l				TALAVIPPFSLSDTLPEVLISLSLI
						YNTLF
25672	56040	Α	25813	215	2160	EINSITSRLSIIPRSPCSASAGWT
						KKAGVPVLASVAAIFLPMWPD
		l				LPIPETTTLPSQCKMILAARTKS
						SPR/RSNIMTHIRIEKGTGGWGG
						PLELKATPGKKIVYITAGTRPAI
						VDKLAQLTGWQAIDGFKEGEP
					1	AEAEIGVAVIDCGGTLRCGIYP
					<u> </u>	KRRIPTINIHSTGKSGPLAQYIVE
						DIYVSGGKEENITVVGDATPQP
				·	İ	SFVGPDYDTSEKLMIVRKKPIG
						VQMLYQVGTVSMLLYTRTIIQT
						KMTPRTIVTISHSVRNAGDSFFS
						IASNQPMPPTAKKIEPLAISLLS
						AEAPTRLSELKARIGEPRWSIEA
						IKPMPISTHKPNMQAITPPARRA
}						ILSPCACESRSRQEPCPPGCHCS
						CPNHSCRSRLKGVLSMEQYLL
						QQKHVGTPPGPFRPPDTNQPCW KTKPSQREEYTRRVGFTGHSNQ
						GAKIIRTHIICRRRTSFALERPGD
						GFQAVVGGTVLMGLPRTKNVG
						LPVTILGRVHRRSSERFPRTSES
						AINRKDAGPKDPVRQRLAYKK
						HPMERRHGPHVVARTFGRAIPT
						LGPHRTVLSTRFQVAKNPLGPI
						GPSPLAVLTWTVPRPKNGRRG
						GKQRARIICQSRREASCNNVCL ARSSGVFKGFSPVSSSGEHTGN
L	<u> </u>	<u> </u>	l			KYSAYSFSLCPLFVFLPLSGRLL

	SEQ ID NO: of peptide sequence		in USSN	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25673	56041	A	25814	11	613	RLSSVFVSKSLASWRSCNCSFS GPLVRLTIRP\LSPPDVRLLLSPS VVHAYNCPHQETQRHHTHHLT P/RLPNHGQIAISAMVYSSPATN/ SMLRQLLIQHVEQIGSVPLVMD AISLVEAHD/RSAAG*SRPLY*R AN*SSRY*VSYARKRVRQAKSR /HGKPSKKVLAKRAVPNRVCRC CNRVKRRCICQPKRRKCMTLPV RATR
25674	56042	В	25815	1	1920	
25675	56043		25816	247	1182	EAVMRQTKTGILLANLGTPDAP TPEAVKRYLKQFLSDRRVVDTS RLLWWPLLRGVILPLRSPRVAK LYASVWMEG\GSPLMVYSRQQ QQALAQRLPEMPVALGMSYGS PSLESAVDELLAEHVDHIVVLP LYPQFSCSTVGAVWDELARILA RKRSIPGISFIRDYADNHDYINA LANSVRASFAKHGEPDLLLLSY HGIPQRYADEGDSLRELYAVA NPNPGPGMGVLLAYMFFGRGS AKQSAGGSGLDGLLRRTPPITP GGYFSVFVWKSLGSWRSCNCS FSGPLVGLTIRPRFTAGRSPTSF AQRCTCL
25676	56044	В	25817	1	651	
25677	56045	A	25818	1	1647	
25678	56046	Α	25819	1	1506	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
<u> </u>	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25679	56047	Α	25820	645	2161	LDRCALNVTAQPSIECQVANAC
						LSPKGEFEYLQIDAPTPQALLSE
						IEKCWHRHRNVWPDHTINLAL
	:					AIHGQVDPVTGVSQTMPQAPW
						TTPVEVKYLLEEKLGIRVMVDN
						DCQLHYCKTAILNWSRKMALS
						ROKFTFERLRRFTLPEGKKOTF
						LWDADVTTLACRATSGAKAFV
ŀ						FQSVYAGKTLRMTIGNINDWKI
	,					DDARAEARRLQTLIDTGIDPRIA
		1				KAVKIAEAESLQAESRKTKVTF
						SVAWEDYLQELRTGISAKTKRP
		1				YSTRYIADHINLSSRGGESPGLG
		l				LLLAFTLFGKGMSKRSAPGAMI
						IHFLGGIHELYFPYVLMKPLTII
						AMIAGGMSGTWMFNLLDGGL
		1				VAGPSPGSIFAYLALTPKGSFLA
						TIAGVTVGTLVSFAITSLILKME
						KTVETESEDEFAQSANAVKAM
		ł				KQEGAFSLSRVKRIAFVCDAG
		ļ				MGSSAMGATT\FRKRLEKAGL
		ł				AIEVKHYAIENVPADADIVVTH
		l				ASLEGRVKRVTDKPLYGPAVV
25680	56048	Α	25821	2	341	
25681	56049	Α	25822	218	469	LICAQGRRSPSRESTRQPKVLTL
						ACHNGSVSVVVLSPNSLYIEVV
						VPS*CNAATGPVVTCLYKGKFQ
				:		PILILLVTSTRVSSCI
25682	56050	A	25823	1	3966	
25683	56051	A	25824	2	728	
25684	56052	Α	25825	528	1123	DAEEGPDHRYLSSAHGGGSAS
		1	İ			QPPSCAVTACRWPWRGRGCRA
						GVWNQPTAAQGRSWPVGKTSC
						*GQGSTCKSQTWGFRGQRLPQ
		l				DAPAPPASLCASFHLVEAGASA
]	l				DSGDTRRMTIR/FASKTVARAK
						RIIRVLRFSPLWVATRSLASTAI
						KIACCASEPGSFLQLWKPRSMA
						PVHRCTWHEACPEGQRQMSCV
						SSTGL
25685	56053	Α	25826	1	492	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25686	56054	Α	25827	1740	3858	ADAAGTEKTATVSRYHFYLRY
						PRSGRSVIDVRSWWRFSIMDAL
						SRSIPRAISICARGKPVVTAKFK
						QGTMPEPSNKAIILPIRFFPGVV
						LMIIKKSGGRWQLSLLASVVIS
1						AFFLNTAYAWQQEYIVDTQPG
						LSTERYTWDSDHQPDYNDILSQ
						RIQSSQRALGLEVNLAEETPVD
:						VTSSMSMGWNFPLYEQVTTGP
			:			VAALHYDGTTTSMYNEFGDST
						TTLTDPLWHASVSTLGWRVDS
						RLGDLRPWAQISYNQQFGENI
						WKAQSGLSRMTATNQNGNWL
						DVTVGADMLLNQNIAAYAALT
						QAENTTNNSDYLYTMGRTRWL
			<u> </u>			TLIGTIITQFALGSVYTWSLFNG
	}					ALSAKLDAPVSQVAFSFGLLSL
						GLAISSSVAGKLQERFGVKRVT
						MASGILLGLGFFLTAHSDNLM
						MLWLSAGVLVGLADGAGYLL
						TLSNCVKWFPERKGLISAFAIGS
				,		YGLGSLGFKF\IDTQLLETVGLE
						KTFVIWGAIALLMIVFGATLMK
						DAPKQEVKTSNGVVEKDYTLA
						ESMRKPQYWMLAVMFLTACM
			:			SGLYVIGVAKDIAQSLAHLDVV
				,		SAANAVTVISIANLSGRLVLGIL
						SDKIARIRVITIGQVISLVGMAA
						LLFAPLNAVTFFAAIACVAFNF
						GGTITVFPSL\ICGSIIASLFGGFY
						VTFYVIFALLILSLALSTTIRQPE
						QKMLHFTYLTTSALFPAGPKRN
25687	56055	A	25828	2	356	
25688	56056	Α	25829	1	1307	MYETLEEVLRSSGDPADQKYV
						ELKARAEKALDDVKKRKLYC
						WRRRQICALRRISRSPYPGHQL
		ĺ				ECIASFDRNFEAMQKEMYGQF
						ENTFMMYLPRLCEHCLNPSCV
						ATCPSGAIYKREEDGIVLIDQDK
						CRGWRLCISGCPYKKIYFNWKS
						GKSEKCIFCYPRIESGQPTVCSE
		Ι,				TCVGRIRYLGVLLYELTAQQFF
						PVWPPDHFVHPF*TLYVLPAPP
						SR*CDRVQLMASLQSVARASRE
						YRWYGFSR*TPRKALALAQLRP
<u></u>						AESSPTAPDCN
		•	•			·_ ·· · · · · · · · · · · · · · · · · ·

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25689	56057	I A	25830	583	1369	RHQLNSIASPAGGRWKDIGWF
23007	30037	``	25050	363	1307	NFPQSNLCLKREKQDESCSLTQ
						ALPSELKVSADNVSLTGAVSLA
		l				SMLTEIFLLQQAQGMPEPGWG
						RITDSHQWNTLLSLHNAQFYLL
						QRTPEVARSRATPLLDLIKTALT
						PHPPQKQAYGGT\LPLPVRFMA
						GPDTNLANLGGALKFNWTLPG
						QPDNTPPGGELVFERWRRLSDN
						SQWIQVSLVFQTLQQMRDKTP
						LSLNTPPGEVKLTLAGCEERNA
						QGMCSLAGFTQIVNEARIPACS
25690	56058	ļ	25831	199	543	QUMCSLAGFIQIVNEARIFACS
25691	56059	A A	25832	1	642	EASIFRRLSVYDNLMAVLQIRD
23091	30039	^	23032	'	042	DLSAEQREDRANELMEEFHIEH
		1				LRDSMGQ*SARPGRCRAGAQ*
		ļ				CV*CR*PRSGQPRQTVRRAE*IS
-		l	1			DSQPVRGQFLHG/VRSPPESDCP
			1			MLSRRCSM*NSSISSFARSSRCS
						ADKSSRICSTAIRLS
25692	56060	A	25833	3	329	ADROGICOTAIRES
25693	56061	A	25834	1974	2195	
25694	56062	A	25835	1	615	
25695	56063	A	25836	2	166	,
25696	56064	A	25837	2	189	STPGNWRFTTTYPLNCATRWK
						M*FLIVATMAPSVYWSLPRNIA
						AAKPTTPPTPSRRSGARGK
25697	56065	Α	25838	64	354	
25698	56066	Α	25839	1123	1845	
25699	56067	A	25840	1120	2097	
25700	56068	Α	25841	124	548	CRQMVGTRRGGEQGKEVIDGŸ
						GKPATFYQMQDNGKPVEGHAS
ļ						QMHYELAKDFVVLTGNAYLQ
1						QVDSNIKGDKITYLVKEQKMQ
						AFSDKGKRVTTVLVPSQLAKA
1						YKG\VAW*KTSA*PSTPGKLSV
		ļ				CWGQTVPVRPPLSTWL
25701	56069	Α	25842	77	514	PRA 4 CL PORVIDA CL 4 POVONA
25702	56070	Α	25843	2487	5738	RRAAGLPGRNDAGLARSHQWA
						GSADTGSEKQHELQLRIQGEPV
						SGQLNLAGSFDRKEERWKG\TL
						SNTRFQTPVGPWSLTRDIALDY
	· .					RNKEQKISIGPHCWLNPNAELC
					:	VPQTIDAGAEGRAVVNLNRFD
			1			LAMLKPFMPETTQASGIFTGKA
			[DVAWDTTKEGLPQGSITLSGRN
						VQVTQTVN\DAALPVAFQTLNL
1						TAELRNNRAELGWTIRLTNNG
						QFDGQVQVTDPQGRRNLGGNV
25700	56051	 	25011	529	222	NIRNFNLAMINPIFTRG
25703	56071	A	25844	538	777	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25704	56072	A	25845	sequence 489	2797	TRRRRTMIALIQRVTRASVTV EGEVTGEIGAGLLVLLGVEKDD DEQKANRLCERVLGYRIFSDAE GKMNLNVHQAAGSVLVVSQFT LAADTERGMRPSFSKGMSQLP GLSRETRESIAMYHLRVPQTEE ELERYYQFRWEMLRKPLHQPK GSERDAWDAMAHHQMVVDEQ GNLVAVGRLYINADNEASIRFM AVHPDVQDKGLGTLMAMTLES VAPYFSGSETGMMTLNRYRLR HMAKQGNRSAKRVEKLLRKPD RLISLVLIGNNLVNILASALGTI VGMRLYGDAGVAIATGVLTFV VLVFAEVLPKTIAALYPEKVAY PSSFLLAPLQILMMPLVWLLNA ITRMLMRMMGIKTDIVDMLLS VLDLEKM/TVDDIMLPRSEIIGI DINDDWKSIL/R/QLSHSPHGAIV LYRDSLDDAISMLRVREAWRL MSEKKEFTKETMLRAADEIYFV PEGTPLSTQLVKFQRNKKKVGL VVNEYGDIQGLVTVEDILEEIV GDFTTSMSPTLAEEVTPQNDGS VIIDGTANVREINKAFNWHLPE DDARTVNGVILEALEEIPVAGT RVRIGEYDIDILDKLSFCFRYSN
						HRRTNRTTIQRITFLHNAQYVT WRNVIRFHHCDGLMHIRVQWD VSFGDHFNAKLTHNIQHRLQRQ LNAFNHRRHIRVSFISHFQRTIQ AINHRQQFVDEFLQREFVGFFNI
25705	56073	A	25846	87	199	QPASVQTAGAGCRL*LHQRQW RECLVLDSKRRGDGVL

	GKSMDKELPWLADNAQ\LELK YKKGKTPLSHRRWPGEPVSVIT
sequence	GKSMDKELPWLADNAQ\LELK YKKGKTPLSHRRWPGEPVSVIT
	YKKGKTPLSHRRWPGEPVSVIT
25706 56074 A 25847 1167 3306	YKKGKTPLSHRRWPGEPVSVIT
23700 30074 A 23847 1107 3300	YKKGKTPLSHRRWPGEPVSVIT
	GSLIQTLGDELLQKAEKKKNIV
	WRYENFSLEWQSAITQAINLIG
	EHKPSIPARTMAALACIAQNDS
	QQLLDEIVQQEGLEYATEVVIA
	RQFIARCYESDPLVVTLQYQDE
	DYGYGYRSETYNEFDLRLRKH
	LSLAEESCWQRCADKLIAALPG
	INKVRRPFIALILPEKPEIANELA
	VPALREFLAATKMPATCTLKGL
	GAVEADYPYYLGMLGMHGTK
	AANFAVQECDLLIAVGARFDD
	RVTGKLNTFAPHASVIHMDIDP
	AEMNKLRQAHLRDEHSWRYD
	HPGDAIYAPLLLKQLSDRKPAD
	CVVTTDVGQHQMWAAQHIAH
	TRPENFITSSGLGTMGFGLPAA
	VGAQVARPNDTVVCISGDGSF
	MMNVQELGTVKRKQLPLKIVL
	LDNQRLGMVRQWQQLFFQER
	YSETTLTDNPDFLMLASAFGIH
	GQHITRKDQVEAALDTMLNSD
	GPYLLHVSIDELENVWPLVPPG
	ASNYKKHHKLKQTFGVKIIPDV
	HDPSQAQPVADVVNVIQLPAFL
	ARQTDLVETHGENRPGQMGNI
	VDKFKEGGNEKVILCDRGANF
	GYDNLVVDMLGFSIMKKVSGN
	SPVIFDVTHALQCRDPFGAASG
	GRRAQVAELARAGMAVGLAG
	LFIEAHPDPEHAKCDGPSALPL
25707 56075 A 25848 1	MATRASVTPMSLFSAGTCLLST
25707 50075 11 25040 1	SCVLSPVGDTAEESCWQRCAD
	KLIAALPGINKVRRPFIALILPEK
	PEIANELVGLECPRTHFHSKEW
	LKVVANDPTAVRKLEHYWSQD
	IFSDREASYMSHENHFGYAACA
	ALLREQGLAAIPRLAMYAHKE DCGSLLVQINHPQVIRTLLLVA
	DKNKPSLQRVAKYHKNFPHAT
	•
	LAALAELLALTEPPA/RPWLSN
05700 56076 4 05040 500	HRRQKAACTAKSTR
25708 56076 A 25849 823 1091	CVRFARINCSDRRSRCFAGFNH
	VLNKALGVHINHFG/YSACAP*
	CRYPPHAAGEFYPDPCRHKGTT
	GYRHRQGYPPPGEPPPRASWLD
	LR
25709 56077 A 25850 714 875	EFSSARISRTSRLKYTNGCCRL
	WSAFCRLRTATTVPPARP*QLR
	PDRSRAQCQ

	equence		09/540,217	4		
25710 5				codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
	6078	Α	25851	2730	2893	WPVGERLAPGDYPVI*TGQLCT
. 1						AAGD/EWRR*KVRRHQPLWRG
						FPNRRYMAWRSA
25711 5	6079	Α	25852	750	929	SFHPAGRRYLYRHSSRSGNITN
						GCCRLWSAFCRLRTATTVPPAR
						P*QLRPDRSRAQCPV
25712 5	6080	A	25853	2	2427	CRQHHRWHSGSTAVYRRVLITS
						KCWNVFIRTKTWTVSILTTSVV
						CASARRVCVPAPRAVSSRCLSV
						TTLIPSASTPW*LAHRISLAAR*
:						AFWFNCRIPAGIDNVKVLERIH
						PDKDVDGFHPYNVGRLCQRAP
						RLRPCTPRGIVTLLERYNIDTFG
						LNAVVIGASNIVGRPMSMELLL
				1		AGCTTTVTHRFTKNLRHHVEN
						ADLLIVAVGKPGFIPGDWIKEG
						AIVIDVGINRLENGKVVGDVVF
						EDAAKRASYITPVPGGVGPMT
						VATLIETRYRQGKMFDGSSIGG
			:			WKGINESDMVLMPDASTAVID
						PFFADSTLIIRCDILEPGTLQGYD RDPRSIAKRAEDYLRSTGIADT
			ļ ,			
						VLFGPEPEFFLFDDIRFGSSISGS
						HVAIDDIEGAWNSSTQYEGGN KGHRPAVKGGYFPVPPVDSAQ
			1			DIRSEMCLVMEQMGLVVEAHH
						HEVATAGQNEVATRFNTMTKK
						ADEIQIYKYVVHNVAHRFGKT
						ATFMPKPMFGDNGSGSPPSQPQ
						SQPTPPEILLQALLSAGCFYAVY
						YVTYKTRLRKRVADVPRRPLFT
						MNRVETNLTWVILMGIALVSV
						GIFFMHNGFLLFRLNSYSQIFSS
1 1						EVSGVALKRFFYFFIPAMLVVY
						FLRQDSKAWLFFLVSTVAFGLL
						TYMIVGGTRANIIIAFAIFLFIGII
						RGWISLWMLAAAGVLGIVGMF
						WLALKRYGMNVSGDEAFYTFL
25713 5	6081	A	25854	502	885	
	56082	A	25855	233	738	SNSCKRTKIISDVTSVFHAIFPAI
		l.,		1		KITAPNSPRLRVNDSATPLTSAG
		1				ESSGSTTLQKLCQRVAPSVSAA
						SSYSGPNLFQHRLHGTHPKRNA
						GKGHSHRNPQPCIGDFDIKLRQ
						PFTVKTLYVKEKGQR*GGDRV
						LPAQRVNPSAHQAACGLGSDS
						GLGSTPSADRKRR

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25715	56083	A	25856	1	2363	MRRQDTGRRGRKRRDAGRWQ
						WRYWGKGSFTPMNATWDVAG
		-			:	KGEWHDSTITLTDLSTGFDQLQ
						YGTMTVEKPRLILDKPIVWVRD
						AQHPSFSGALSLDAGQTLFTGG
						SVLPPSTLKFSVDGRDPTYFLFK
						GDLHAGEIGPVRVNGRWDGIR
						LRGNAWWPKQSLTVFQPLVPP
						DWKMNLRDGELYAQVAFSAA
						PEQGFRAGGHGVLKGGSAWM
						PDNQVNGVDFVLPFRFADGAW
						HLGTRGPVTLRIAKQRKEMRN
						NNLRQFKLLGPDVGFDSINDRP
						MAEELSKLLSKQNEENLLPKTI
						LYCLNPRDNEVLGTMIENGASF
						AANPLYFDPKNIVELAIEAGCN
						CVASTYGVLASVSRRYAHRIPF
İ			<u> </u>			LVKLNHNETLSYPNTYDQTLY
						ASVEQAFNMGAVAVGATIYFG
						SEESRRQIEEISAAFERAHELGM
				1		DGGWIGLMIKPLGRWSLIMEID
						EGFAVGMSPAELSAEQLLSKL
						WLWEGKAERYGWGRKSMDKE
						LPRLADNAQLELKYKKGKTPLS
						HRRWPGEPVSVITGSLIQTLGD
						ELLQKAEKKKNIVWRYENFSLE
					}	WQSAITQAINLIGEHKPSIPART
						MAALACIAQNDSQQLLDEIVQ
						QEGLEYATEVVIARQFIARCYE
						SDPLVVTLQYQDEDYGYGYRS
		l				ETYNEFDLRLRKHLSLAEESCW
		ŀ				QRCADKLIAALPGINKVRRPFIA
25716	56084	Α	25857	1	4440	MNRFDGAKQNVVGADAHHLA
		ŀ				DTAVKRNQSLFQLRASGHPRLP
						AGGGKAVFHGRFAAKKIEAVA
						FYFRHLIALLVVQMHFKLVKG
						NVDVVFTAQLIDSTVQLIDSTTT
						VTQFTHIQTQLMRSVEWLTDN
						GSCYRANETRQFARMLGLEPK
						NTAVRSPESNGIAESFVKTIKRD
						YISIMPKPDGLTAAKNLAEAFE
1						HYNEWHPHKAGTLTNEQWQQ
						VTAELHDRMMETVFFALDDAE
						QLFAHHQPTPVTSVDLLGQG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
] 				sequence		
25717	56085	A	25858	1	3705	MEEYGGVDVRLYRIPDPMAFL
						RQQKNLHRIVVQPQYLGDGLN
						NTLTWLWDNWYGKSRRVMQR
						TFSSQSRQNVTQALPELQLGNA
						IIKPSRYVQNNQFSPLKKYPLVK
						QFRYPLWQAKPFEPQQGSRGL
						YLVEAMVGGYRATTGGVVSDT
						VALSKVSSKELLVWTAGKKQV
				•		NEPGSEIFGLTALRITAENQDCA
						SLTPETFLPRIHVIKGVNISTATV
						CRQCEDAPCANVCPNGAISRDK
		·				GFVHCEARTLHLFAKT
25718	56086	Α	25859	123	923	TDSPYERGHLMSNLLGPRDAN
	1					GIPVPMTVDESIASMKASLLKKI
						KRSAYVYRVDCGGCNGCEIEIF
						ATLSPLFDAERFGIKVVPSPRHA
						DILLFTGAVTRAMRSPALRAW
						QSAPDPKICIS\YGACGNSGGIF
			•			HDLYCVWGGTDKIVPVDVYIP
						GCPPTP\AATLYGFAMALGLLE
						QKIHARGPGELDEQPAEILHGD
						MVQPLRVKVDREARRLAGYRY
						GRQ\VADDSFTQLGQGEEQVAR
						WLEAENDPRLNEIVSHLNHVVE
25719	56087	A	25860	1337	1516	KLKSKSAKRHKFITPILAWRILI
23,17	30007	`	23000	1337	1310	HYLQRVAV*WGSRLL*LFLITIP
						SRWRLLPHNTGKG
25720	56088	Α	25861	1	545	MARLAALKACETVYSMGVRSH
25,20		`		1		RRSPLEHLWQLKDQLVNEGEL
						VLETLVIDGDENTVLVPGDRYA
						QMRNVYFIPSALALKNWLKKC
			:			GFVDIRIADVSR/ITTTEEQRRTE
						WMVTESLADFLDPHDPGKTVE
						GYPAPKRARNRNRAEDHRORL
						RHRTRKKQTHGQRRRHRPEPA
						QRQPQTGTTD
25721	56089	A	25862	620	1117	νηνι ν 1011 μ
25722	56090	Α	25863	1	1188	
25723	56091	A	25864	231	407	
25724	56092	A	25865	1	501	
25725	56093	В	25866	1	2373	
25726	56094	Α	25867	1	2211	

SEQ ID NO:	SEQ ID NO: of peptide sequence	t .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25727	56095	A	25868	2093	3589	GGINGKNFWFFLSPFTPCVGRC HSRVQLANSHLVDVDETPPSSA AVHLYATDADSIGNCGTDREFC PCNPIRRHAFCCERLQNGYHRS GYAFHLGHYGDRKVITNFYSLI AKNHLSHWLETLPAQIANWQR EQQHGLFKQWSNAVEFLPEIKP YRLDLLHSVTAESEEPLSAGQI KRIETLMRNLMPWRKGPFSLY GVNIDTEWRSDWKWDRVLPHL SDLTGRTILDVGCGSGYHMWR MIGAGAHLAVGIDPTQLFLCQF EAVRKLLGNDQRAHLLPL/GIE QLPALKAFDTVFSMGVLYHRR SPLEHLWQLKDQLVNEGELVL ETLVIDGDENTVLVPGDRYAQ MRNVYFIPSALALKNWLKKCG FVDIRIADVSVTTTEEQRRTEW MVTESLADFLDPHDPAIFQIWP CTFMHAYHAFFDPVSHHRACQ NLTTFVPDTHQIAIRNTKACSIF VSFALRTTHSCAISHSSLEDPPAI
25728	56096	A	25869	231	1785	LFKIADRMEQNLELLATAGNLG DDEVHRYSELRTRTVPSYPPER SGEAKTAAQRAGTAFRRGRGA GVPFNLQQRSDRWCS*RSQQN A*RLIGLHGQNSRSVPQLPMES ASVA*RCGSIIEKTTLSSRPFYW CPGCQHYADRFGA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25729	56097	A	25870	1818	3311	RMAIQFRTLSLRWCWKISLVSS FTVVKAMTIRIVAMRQLLRQQ KQRTFVC*SS/TVDGVLSDGLIY MGNNGEELKAFNVRDGYGIRC ALTSDIEVAIITGRKAKLVEDR\ LPHWGSLTCIRGSQTN*SPLAIC WKNWRLP/QENVAYVGDDLID WPVMEKVGLSVAVDDAHPLL GTEIDAYIENADYQYNKKRLRE IDRRVRYLTKCLENLKIVDYSP QQEGKVFFGAWVEIENDDGVT HRFRIVGYDEIFGRKDYISIDSP MARALLKKEVGDLAVVNTPAG EASWYVNAIEYVKPRLNQTVR LLLEHEMGQVWISGEISNFTQP ASGHWYFTLKDDTAQVRCAM FRNSNRRVTFRPQHGQQVLVR ANITLYEPRGDYQIIVESMQPA GEGLLQQKYEQLKAKLQAEGL FDQQYKKPLPSPAHCVGVITSK TGAALHDILHVLKRRDPSLPVII YPAAVQGDDAPGQIVRAIELAN QRNECDVLRPCVWCRRVRCRR
25730	56098	A	25871	1	3465	QRIVEED VERI EV WERR VRERR
25731	56099	A	25872	1	3126	
25732	56100	A	25873		1722	MSSLSQAASSVEKRTNARYWI VVMLFIVTSFNYGDRATLSIAG SEMAKHIGLDPVGMGYVFSAF SWAYVIGQIPGFVDIFSGFGIIV ALFTLRFLVGLAEAPSFPGQSRI VGDWFPAQDRGTGVSIFTPFIL RKGDLRPIWVVILMAGQLHAL MSTLEDTPFSNKAGMRLRRWI NAAVMRLNPLACLATRSNQVN WLLLTGNISVYPVGSSEPTRR STPINSTPHFNKTILPINARYSPT LPSSDGTRWTTRIDDKPTVGHK FARYWDVELREIP\MRPG\QLFR DPKRMIEGCAENPL\GV\VPTFG VTYAGNYEFPQPLPDALNKFQ ADPGIDIDMHIDAASGGFLAPF VAPDIVWDFRLPRVKSISASGH KFGLAPLGCGWVIWRDEEALP QELVFNVDYLGGQIGTFAINFS RPAGQVIAQYYEFLRLGREGYT KSTQPARGSPPANNCQAQPETP DRAVIHPTDGYESAATIIVLSPP TASDSDPAHRDQRVVKTDAWS VVPGSTSEYPRRYKAEWFCQPF AVLPAILAWSAALWRYQPRAR RRKADYGCRALNSLKNELTTP

SEQ ID NO:	SEQ ID NO: of peptide sequence	I .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25733	56101	Α	25874	1	2385	
25734	56102	Α	25875	1	1767	
25735	56103	Α	25876	1	2079	
25736	56104	A	25877	368	1261	TSPRARPAAQGSRGAFL/HLTE KGVNRFAFYGLPESSGKRWAT EREYAFRQLVAEEKYRGVVYQ GLETAPENWQHAQNRLADWL QTLPPQTGIIAVTDARARHILQV CEHLHIPVPEKLCVIGIDNEELT RYLSRVALSSVAQGARQMGYQ AAKLLHRLLDKEEMPLQRILVP PVRVIERRSTDYRSLTDPAVIQA MHYIRNHACKGIKVDQVLDAV GISRSNLEKRFKEEVGETIHAMI HAEKLEKARSLLISTTLSINEISQ MCGYPSLQYFYSVFKKAYDTT
25737	56105	A	25878		868	PKEVFAIVNSEVML MTSLKNSMKWDEERFGLEYDL DIYMIVAVDFFNMGAMENKGL NIFNSKYVLARTDTATDKDYLD IERVIGHEYFHNWTGNRVTCRD WFQLSLKEGLTVFRDQEFSSDL GSRAVNRINNVRTMRGLQFAE DASPMAHPIRPDMVIEMNNFYT LTVYEKGAEVIGM\TPPLFGQE NFQNVILTCALEDPTRKVVLKP LCRPPLSLCNRRKRVMCRIFTSS ASTLRLSRRCAVCWNSRITVSM RSSRRVTSVWLSAPTPIILSPAIF IVRWWLLDSNPLIYYKAWSCW CSRK
25738	56106	Α	25879	2097	2512	NTPVRPDVGT*TEEHGGAESGE *RNSRELRENDKA*LHQYHAQP DGKNLAEAFEHYNEWHPHSAL GYWRTVD/DMRGR*PQEPSLMI RAGRSSSPVQWQRIPLLIPGRKG DRMQKTVIPGSLFAGMPYAQPS IKTELAG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			:	sequence		
25739	56107	Α	25880	71	2613	QRKPTSSRKGLSARMMSGLTG
						TITTVSTGWCTTRGHNMKSRW
		l				SKVAIFTTR/SGRRVAKRVWRR
		1				ERDLTGWMSLSRKPQVTWYG
			,			WDGDRLTTIQNDRSRIQTIYQP
		1				GSFTPLIRVETATAVRRRRRWQ
		1	ŀ			WGIPAGTVQMLDRLESEILADR
		İ				VSEESRRWLASCGLTVEQMKK
ļ		1				TSIHASIVRHKLLTAEQALKLLG
		1				EMFVYHMPFNRALGMELERYE
			,			KEFAQLAFKNQPMMVGNWAQ
			,			SILHGGVIASALDVAAGLVCVG
		İ				STLTRHETISEDELRQRLSRMGT
						IDLRVDYLRPGRGERFTATSSLL
					•	RAGNKVAVARVELHNEEQLYI
						ASATATYMSKSMFLAVATILIS
						KSSLSRLRRPSTIVCGIPISSISAN
						ITPGRSLRSSSSTSKTLCTQFAV
						QFLSKLLYAEGFVHVHRQNRH
						LERRDSVRPYDASFVVVLLNRR
	ļ	i				SYHTRYPDTVATHRQDLVTAIF
İ						TLYGGFQCFGVLGTQLEDVTNF
						DTTFDQQPTLAIRA WIASHHVT
		ŀ				DICNFRGSDIAIPVDAEVVFTID
						VSTSSEITHRRNVDLFPDQLTVT
						RSQTFTAYERLSTALTVAQVCG
						VQRLCNYYSARLTPLPGPDSTR
						ESNHRLAQITQYARQLASSPSII
						DNRSRQHLNDVGLTAWDCVIIS
						QIIGFIGFQARTIATFQAYLGHP
						VRWLPGLEIQNYADASLFADES
						LRWRSSYEVEKLPEEHTKSSTA
25740	56108	Α	25881	71	3478	CCLSSHPSTTATALRSLSPVRK
						WPKISALIPWEWAMCSLLSHGL
		ŀ				MLSGRSLVAWCWTVLVQNAS
						TSGRSLSGRCLPCCKASSISLVD
						SALSFLLFTLRFLVGLAEAPSFP
						GNSRIVAAWFPARKGTGIINE\IS
						P\LANTALEGLGGSGDALCTQC
						EAEGFRHITYYLDRPDVLARFT
						TKIIADKIKYPFLLSNGNRVAQG
					1	ELENGRHWVQWQDPFPKPCYL
						FALVAGDFDVLRDTFTTRSGRE
						VALELYVDRGN

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25741	56109	A	25882	90	629	PGDWGRRQLIGS*KTPGSVFKK GKSSTPN\DPFLKDKTQDRSGL ATFTMDGQI\RLFYQ\DISGNHS GK\QSLPPAQVIVSNSVDHLKIT GVEDHKPIFAGAGKPYQNLRQF IDEGNYTSGDNHTLRDPHYVED KAHKYLV/FEANTGTENGYQG EESLFNKAYYGGGTNFFRGSSR VDLQAC
25742	56110	Α	25883	1	3066	
25743	56111	Α	25884	104	547	LFPRLLSCLTTPPHCSFSICFVIC SRTLILKGSSLMYVFCLPNTAIV MALSPRGWRSKFGMPVDSKGP PWLFKFLKNGLNFLLHSVGTRD LSRLSTILSPLYNTVAPGAN*CR ELKLVLDADDVLSTEVKRVITS SASNITPAFFCS
25744	56112	Α	25885	3	4288	
25745	56113	A	25886		2222	DFADFGTTIKQDFRLLGQTSVD RLLQLSQGQAVKGNQLLPVSLP QHTIPDVFIWMLSNNRRVAYA RIASKDLLYSPVAGQMGKHCG KIKTHFLKIPKQQQNEKYQVPQ FDQSTIKNIESAKGLDVWNSWP LQNADGTVAEYNGYHVVFALA GSPKDADDTSIYMFYQKDFRLL GQTSVDRLLQLSQGQAVKGNQ LLPVSLVKRKTTLAPNTQTASP RALADSLM\SWHDRFPDWKAG RILPISEPPSNRIFACWGKPAWT ACCNSLRARR*RAISCCPSHW* KEKPPWRPIRKPPLPARWPIH*C SWHDRFPDWKAGRKGSDAHRF GSWKFQLQRVSG\SATFTWGG KIR\LYSADYSGNHYGKQSLPS GQVNGSNSDDTLKINGVEDHK TILDGDGKTYQNVQQFIDEGNY TSGDNHTLRDPHYVEDKGHKY LVFEANTGTENGYQGEESLFNK AYYGGGTNFFRKESQKLQQSA KKRDAELANGALGIIELNNDYT LKKVMKPLITSNTIRKSDGTLQ EHDGICEIHVAKYAEIFGLTSAE ASKDIRQALKSFAGKEVVFYRP EEDAGDEKGYESFPWFIKRAHS PSRGLYSVHINPYLIPFFIGLQNR FTQFRLSETKEITNPYAMRLYES LCQYRKPDGSGIVSLKIDWIIER YQLPQSYQRMPDFRRRFLQVC
	1	1	0.500-	ļ.	2612	VNEINSRTPMRLSYRMGVRGK
25746	56114	В	25887	1	2640	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25747	56115	Α	25888	1291	1629	
25748	56115	A	25888 25889	1187	2925	NSLQRSHPNAANS/WHNKLWR SHSDSDLSDHHEPICKPGLELN KKDITTSADQIAEVKTMESHPPI PPVFVEHMVPQDANQKGLCTK ERMICLEFTSREFHAGQIEDELN LNDINGCSSGSTLEDPQFNDDSP EARKITRRWRIGEAADLVGVSS QAIRDAEKAGRLPHPDMEIRGR VEQRVGYTIEQINHMRDVFGTR LRRAEDVFPPVIGVAAHKGGV YKTSVSVHLAQDLALKGLRVL LVEGNDPQGTASMYHGWVPD LHIHAEDTLLPFYLGEKDDVTY AIKPTCWPGLDIIPSCLALHRIET
						ELMGKFDEGKLPTDPHLMLRL AIETVAHDYDVIVIDSAPNLEDP RVGVMARGNAITLPVCGRDVK FTLEVLRGDSVEKTSRVWSGNE RDQELLTEDALDDLIPSFLLTGQ QTPAFGRRVSGVIEIADGSRRR KAAALTESDYRVLVGELDDEQ MAALSRLGNDYRPTSAYERGQ RYASRLQNEFAGNISALADAEN ISRKIITRCINTAKLPKSVVALFS HPGELSARSGDALQKAFTDKEE LLKQQASNLHEQKKAGVIFEAE EVITLLTSVL
25749	56117	В	25890	1	1954	
25750	56118	A	25891	1	454	MERRNRRTGRTEKARIWEVTD RTVRTWIGEAVAAAAADGVTF SVPVTPHTFRHSYAMHMLYAG IPLKVLQSLMGHKSISSTEVYTK VFALDVAARHRAIR\VPRQQGD YRTRIWKFEDGLSNVLVIQLNK LIICVMCLVRDCDVLKTYFHL
25751	56119	A	25892	10	144	FLFIGKGFLKGIT*PGQIVQRG*E KKSNT*FRQFGGVDTAGNNLT

SEQ ID SEQ NO: of pe seque	otide hod	09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25752 5612	20 A	25893	888	892	TVDLDACTKGVLTFTTALLAG GANQAFAKENTKKAYKETYGV SHITRHDMLQIFKQQQNEKYQV PQFDQSTIKNIESAKGLDVWDS WPLQNADGTVAEYNGYHVVF ALAGSPKDADDTSIYMFYQKV GDNSIDSWKNAGRVFKDSDKF DANDPILKDQTQEWSGSATFTS DGKIRLFYTDYSGKHYGKQSLT TAQVNVSKSDDTLKINGVEDH KTIFDGDGKTYQNVQQFIDEGN YTSGDNHTLRDPHYV/EDKGH KYLVFEANTGTENGYQGEESLF NKAYYGGGTNFFRKESQKLQQ SAKKRDAELANGALGIIELNND YTLKKVMKPLITSNTVTDEIER ANVFKMNGKWYLFTDSRGSK MTIDDLWKFELENDDLEYLVE ENSKQKTIQDVIWLFLTAYTHI RGATQAFAKENNQKAYKETYG VSHITRHDMLQIPKQQQNEKYQ VPQFDQSTIKNIESAKGLDVWD SWPLQNADGTVAEYNGYHVVF ALAGSPKDADDTSIYMFYQKV GDNSIDSWKNAGRVFKDSDKF DANDPILKDQTQEWSGSATFTS DGKIRLFYTDYSGKHYGKQSLT TAQVNVSKSDDTLKINGVEDH KTIFDGDGKTYQNVQQFIDEGN YTPGDNHTLRDPHYVKTKAMR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	1	
25753	56121	Α	25894	1424	3807	HPLWKWLEGDMNMNIKKIVK
23733	30121	^	23034	1424	3007	i i
		ŀ				QATVLTFTTA/LLAGGATQAFA
İ						KENNQKAYKETYGVSHITRHD
						MLQIPKQQQNEKYQVPQFDQS
						TIKNIESAKGLDVWDSWPLQN
		l				ADGTVAEYNGYHVVFALAGSP
		1				KDADDTSIYMFYQKVGDNSIDS
		l				WKNAGRVFKDSDKFDANDPIL
						KDQTQEWSGSATFTSDGKIRLF
						YTDYSGKHYGKQSLTTAQVNV
						SKSDDTLKINGVEDHKTIFDGD
						GKTYQNVQQFIDEGNYTGDPL
		1				EAETAVINHKKRKNSPRIVQSN
		1				DLTEAAYSLSRDQKRMLYLFV
		1				DQIRKSDGTLQEHDGICEIHVA
						KYAEIFGLTSAEASKDIRQALKS
						FAGKEVVFYRPEEDAGDEKGY
		1				ESFPWFIKRAHSPSRGLYSVHIN
						PYLIPFFIGLQNRFTQFRLSETKE
	ļ					ITNPYAMRLYESLCQYRKPDGS
		1				GIVSLKIDWIIERYQLPQSYQRT
						PDFRRRFLQVCVNEINGAVIGIP
						CVSIRKPDGSGIVSIKIAWIIERY
		1				QPPQSYQRMPDFRRRFLQSRPA
1						CMHDWLCAEALAWSIQTASYL
		1				VTMQVNLTSLSSDTDRDLSVVS
		1				NSGWVSSGSLVRFNTIKTSSGEI
		1				KRTVPRILPDPDDPRSAIAEAPS
		İ				EMPGHEVPVEEHFPEAGTNSGS
						PQGARKGDESMTKASDSSSPSC
		l				SSGPRVPKGAAPGSQTGKKQQS
						TALOASTLAPANLLPKAVHLA
25754	56122	A	25895	3	2356	
25755	56123	Α	25896	1	2610	
25756	56124	Α	25897	1	2496	· · · · · · · · · · · · · · · · · · ·
25757	56125	В	25898	1	3198	
25758	56126	Α	25899	1	5274	
25759	56127	Α	25900	1	585	
25760	56128	A	25901	[]	2469	
25761 25762	56129	A	25902	1	1914	
	56130	A	25903	1267	2259	VI VTI IIMDI QWDINIQQVQQQQ
25763	56131	Α	25904	1367	1959	KLVTLHMDLSWR\NSGYQSQE
		1				T*K*PANRPVKRPH*GGI*\FSRD
						QKRMLYLFVDQIRKSDGTLQE
						HDGICEIHVAKYAEIFGLTSAEA
		1				SKDIRQALKSF/DGEGSIYSR*A
		1				HGAA*PSHRKDRESADLGSDG
						QNDKGH\QYLVFEANTGTENG
						YQGEESLFNKAYYGGGTNFFR
						KESHKLQQSAKKLDAELSNGA
						LGFIE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25764	56132	Α	25905	1	2418	
25765	56133	A	25906	3	1918	
25766	56134	Α	25907	1242	1463	
25767	56135	A	25908	1907	5097	TSKKIVKQAPVLTFTTA/LLAGG
						AIQAFAKENNHKAYKETYGVS
						HITRHDMLQIPKQQQNEKYQVP
						QFDQSTIKNİESAKĞLDVWDSW
						PLQNADGTVAEYNGYHVVFAL
						AGSPKDADDTSIYMFYQKVGD
						NSIDSWKNAGRVFKDSDKFDA
						NDPILKDQTQEWSGSATFTSDG
						KIRLFYTDYSGKHYGKQSLTTA
						QVNVSKSDDTLKINGVEDHKTI
						FDGDGKTYQNVQQFIDEGNEGI
						LPISEPPIKODFRLLG
25768	56136	Α	25909	610	2303	SLPNLDNAAICSSSSSPTRTR*SL
						SEGATQ\AFAKEKYPHKHTKKR
						SGVFHITRHDMLQIPKQQQNEK
						YQVPQFDQSTIKNIESAKALDV
						WDSWPLQNADGTVAEYNGYH
						VVFALAGSPKDADDTSIYMFY
						QKVGDNSIDSWKNAGRVFKDS
	i					DKFDANDPILKDQTQEWSGSA
						TFTSDGKIRLFYTDYSGKHYGK
						QSLTTAQVNVSKSDDTLKINGV
						EDHKTIFDGDGKTYQNVQQFID
						EGNYTSGDNHTLRDPHYVEDK
			,			GHKYRGPLESPSTHQAEFNPTS
						CVSSLGTLQGFPAPAWLALAHP
						VHPLKHKSGGSNRLSAAIWGIK
						RKPARVCPGTGIHASSQIQGEW
						RTECAVGPKAKAKATAGWRR
						GNNQHISSTYDINRADTQVRRA
						VNNYDIIVMSNSFNGQSEHQTY
						ESIVIDSAPNLGIGTINVVCAAD
						VLIVPTPAELFDYTSALQFFDM
						LRDLLKNVDLKGFEPDVRILLT
						KYSNSNGSQSPWMEEQIRDAW
						GSMVLKNVVRETDEVGKGQIR
						MRTVFEQAIDQRSSTGAWRNA
						LSIWEPVCNEIFDRLIKPRWEIR
25760	56137	Δ	25910	1	2103	DOTALI VOILER DREIRI RWEIR
25769	56137	A	25910	1	2103	LSIWEFVCHEIFDRLIKFKWEIF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25770	56138	A	25911	1	2124	MNMNIKKIVKQATVLTFTTALL
						AGGATQAFAKENNQKANKETY
						GVSHITRHDMLQIPKQQQNEKY
		l				QVPQFDQSTIKNIESAKGLDVW
		1				DSWPLQNADGTVAEYNGYHV
						VFALAGSPKDADDTSIYMFFKR
		ŀ				GAIFRVHKHAVNPMSPKCRRPG
						GRQAYPLVNWEDRNGRSQKTV
		l				HTEGDMNMNIKKIVKQATVLT
		ŀ				FTTALLAGGATQAFAKENNQK
		ŀ				AYKET/YPKQQQNEKYQVPQF
•		ŀ				DQSTIKNIESAKG\LDVWDSWP
		l				LQNADGTVAEYNGYHVVSALA
1						GSPKDADDTSIYMFYQKVGDN
						SIDSWKNAGRVFKDSDKFDAN
						DPILKDQTQEWSGSATFTSDGR
						RSLESTTTAARPIWRKDVGGDQ
						TQEWSGSAPFTSDGKIRLFYTD
						YSGKHYGKQSLTTAQVNVSKS
						DDTLKINGVEDHKTIFDGDGKT
						YQNVQQFIDEGNYTSGDNHTL
						RDPHYVEDKGHKYLVFEANTG
						TENGYQGEESLFNKAYYGGGT
						NFFRKESQKLQQSAKKRDAEL
						ANGALGIIELNNDYTLKKVMKP
						LITSNTVTDEIERANVFKMNGK
						WYLFTDSRGSKMTIDGINSNDI
		İ				YMLGYVSNSLTGPYKPLNTTG
						LVLQMGLDPNDVTWASLEPHE
						SFQWVRGLASSGVKLQTSVVL
						QLIKAMWTQRVSSSKVYCKEQ
						MNNASTMSKRTSAGCHCWQG
25771	56139	A	25912	1	3987	
25772	56140	Α	25913	1	2235	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25773	56141	A	25914	1	1950	MNMNIKKIVKQATVLTFTTALL
						AGGATQAFAKENNQKAYKETY
ł						GVSHITRHDMLQIPKQQQNEKY
						QVPQFDQSTIKNIESAKGLDVW
						DSWPLQNADGTVAEYNGYHV
						VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
			:			KFDANDPILKDQTQEWSGSATF
						TSDGKIRLFYTGSLNSSKTEKY
						QVPHIDQSTIKNIESAKGLDVW
						DSWPLQNADGTVAEYNGYHV
						VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
			:			KFDANDPILKDQTQEWSGSATF
						TSDGKIRLFYTDYSGKHYGKQS
						LTTAQVNVSKSDDTLKINGVED
						HKTIFDGDGKTYQNVQQFIDEG
						NYTSGDNHTLRDPHYVEDKGH
						KYL/VFEANTGTEEHPQPQ\ERP
						RTQSFTSAFAERRECIPNVPADT
						KLSKIKTLRLATSYIAYLMDLL
						AKDDQNGEAEAFKAEIKKTDV
					İ	KEEKRKKELASKCLDLEQLGAS
			<u> </u>			VEPTGNLRTKITKEKPRHTGPPE
						VVVPGCCPHRSRAYKSDKYAH
						TLTVTASQHAPPPPTHMEGFEL
						FHLPDLCSPSQDAQTTGRTQMK
						PDHSPRPSHRVPQAKGNNVVIT
						SYMTNRGFFEDKKATFAPSFLM
25774	56142	A	25915	1	2448	NIKGNKTSVVKNSILEQGQLTV
25775	56143	A	25916	1312	1974	
25776	56144	A	25917	2	1778	
25777	56145	A	25918	1366	2673	
25778	56146	В	25919	1	1938	
25779	56147	В	25920	1	2262	
25780	56148	Α	25921	1	2100	
25781	56149	В	25922	7145	7154	
25782	56150	Α	25923	2085	11232	VNSEGSEPADRKKPVHTEEAM
		1	ł			NMTIKKIVKQATVLTLTTALLA
						EGATQAFAKENNQKAYKETYG
						VSHITRHDMLQIPKQQQNEKYQ
						VPQFDQSTIKNIESAKGLDVWD
						SWPLQNADGTVAEYNGYHVVF
						ALAGSPKDADDTSIYMFYQKV
						GDNSIDSWKNAGRVFKDSDKF
						DANDPILKDQTQEWSGSATFTS
						DGKIRLFYTDYSGKHYGKQSLT
						TAQVNVSKSDDTLK\INGVEDH
0.5700	56151	 	0.500.4	1.52	202	KPILDGGGKTYQNVQQFI
25783	56151	Α	25924	153	293	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25784	56152	Α	25925	420	687	
25785	56153	A	25926	242	491	WLHVQPLHDEHHCWI*YPADT DGVRPAFQRSPALSPEHGCVPY TPARQGSVRSKGPCTSALLYVE CSIAAPDRKHSCHKSDQS
25786	56154	Α	25927	2	113	
25787	56155	A	25928	779	862	NSTLPRCASTTFARY*AIPIKVFS ILP
25788	56156	A	25929	282	443	FSAGFLNLRYAQKFPSSYPPV** CYRPIYYPASDPCRNRLLYPAD VPYQALLD
25789	56157	Α	25930	3	89	
25790	56158	Α	25931	614	823	
25791	56159	Α	25932	1229	1427	
25792	56160	A	25933	1965	2682	FAVPCRTSFGSIETVSHPGRDT* SA*AYRVQVDSHRPRLAHNVQ HRT*RGEVLLAGVPRHVAEREI ATLAGSFSLHEQNIHNLPRDQG PGNTVSLEVESENITERFFVVGE KRVSAEVVAAQLVKEVKRYLA STAAVGEYLADQLVLPMALAG AGEFTVAHPSCHLLTNIAVVER FLPVRFSLIETDGVTRQLLGVSY RILAMGHAEFLIQIADMRNDGG WRDFQFSGNLVMDEPNRSAQT YIKLVKSRLGTTKR
25793	56161	Α	25934	261	563	RHSRGRSALRQISPEC*IHVCSPP PVQQLHPGYGNAIPLSYGHRNQ PDESDDAPVPPGWCQRYNLHQ KSTAHCGGYTIHADLVKTRCW LAPHRLPPGPGR
25794	56162	В	25935	1	1329	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ł		sequence		
25795	56163	A	25936	761	3224	I RGGAVTAKAPPDISAISALTAV
23173	30103	ľ`	23730	, , ,		KHGNCSSLTPLLNPPGSDVIVC
						AEMDEQWGYVGAKSRQRWLF
		l				YAYDSLRKTVVAHVFGERTMA
						TLGRLMSLLSPFDVVIWMTDG
						WPLYESRLKGKLHSNEGDAVL
						LEIEQEVSSVDSSFNSTLGPTFN
		:				TRTIQNAVLVKTGETVVLGGLL
						DDFSKEQVSKVPLLGDIPLVGQ
						LFRYTSTERAKRNLMVFIRPTII
						RDDDVYRSLSKEKYTRYRQEQ
						QQRIDGKSKALVGSEDLPVLDE
			•			NTFNSHAPAPSSRWALAQRIGV
						SRVDMQQHVRRPPVNLSPKRE
						NVIAGYLSNSVARKQRHIQTGT
						VFHQLMNLLRDGINIIFMNFKY
						QFIVYLHFLLSPKWVMYNRKC
						VNGMLLIASVPATWMLSRYRS
						SHRWIAKEIPMGKTQPLPILLLG
		1				/GGRRIGLALA\WHFIIQKQPVIV
						SYRTHYPAIDGLINAGAQCIQA
						DFSTTDGVMAFADELLKSTHG
						LRAILHHASAWMAEKPGAPLA
						DVLACMMQIHVNTPYLLNHAL
						ERLLRGHGHAASDIIHFTDYVV
						ERGSDKHIAYAASKAALDNMT
						RSFARKLAPEVKRQTATRAGH
					İ	RGINQLSRYYRRALARQNQRR
						MVELRSLRFMYRHRPRSFVLW
					İ	QATRLHRLNIAVRHRKPDARPL
						PAIQRNTDIAIKQPKRAVITGNH
						HRSTFVPTRLAQTGSEIRRHAM
25796	56164	Α	25937	3	543	
25797	56165	Α	25938	209	343	LSPPMSSFLATVFRSIAWLLTAK
						SIRYRTT*RWIRCAAVWLTMD
25700	56166	<u> </u>	25020	547	640	
25798 25799	56166 56167	A	25939 25940	174	583	
25800	56168	A	25941	2001	2610	DVLV\RKHNGHWTVELYSDSIP
23000	20100		22771	2001		RLQINQHYASMCNNARNDGDS
					1	QFIRSNLQDAKWLIKSLESRND
					1	TLLRVSRCIVEQQQAFFEQGEE
		1				YMKPMVLADIAQAVEMHESTI
		1				SRVTTQKYLHSPRGIFELKYFFS
					1	<u> </u>
		1				SHVNTEGGGEASSTAIRALVKK
						LIAAENPAKPLSDSKLTSLLSEQ
		1				GIMVARRTVAKYRESLSIPPSN
	<u> </u>	1				QRKQLV
25801	56169	Α	25942	329	484	TRQLSLLVRMTL*AFVRKRRSS
		İ				LA*KPLWSWLSSVKSQPGWKL
		<u></u>	<u></u>		<u> </u>	MLARWKSF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25002	56170		25042	11	1785	
25802 25803	56170	A	25943 25944	2493	2696	
	56171	A		<u> </u>	1861	
25804	56172	A	25945 25946	1	700	
25805	56173 56174	A A	25947	456	2413	MPVIKVRENEPFDVALRRFKRS
25806	30174	A	23947	1	2413	CEKAGVLAEVRRREFYEKPTTE
		1				RSRYDLVDRNNNIVLEYRKKEL
						VRLTLTDPVTGKSGEVKSLVSS
						LQTKYALKGYNVEATALEAAG
}						GKVVTTGKDILVRKPTLDLPLE
						VRRKMWFKPFMQSYLVVFIGY
						LTMYLIRKNFNIAQNDMISTYG
						LSMTQLGMIGLGFSITYGVGKT
						LVSYYADGKNTKQFLPFMLILS
						AICMLGFSASMGSGSVSLFLMI
						AFYALSGFFQ\ST\GGSCSYST\IT
						\KWTPRRKRGTFLGFWNISHNL
						GGAGAAGVALFGANYLFDGH
				1		VIGMFIFPSIIALIVGFIGLRYGS
						DSPESYGLGKAEELFGEEISEED
						KETESTDMTKWQIFVEYVLKN
						KVIWLLCFANIFL\YVVRIGIDQ
						WSTVYAFQELKLSKAVAIQGFT
						LFEAGALVGNGCSYAQGMWG
						KTAETSDLQDLLIAALQGLSAW
ŀ						AVKAREYGIINHDVDSFAPRAF
						FSTLTNVNFDSPRIVGYAREAIA
	1					LREALKAQCLAVDANARVDNP
1						MADLQLVSDDLGELQRQAAEF
						TPNKDKAAIGENILGLRLLCLY
						GLKVFNGREDQLDSCYHGNRQ
						LRGSLRLLFLAFIRQRLRDLIITA
						HRVIRVNRQHAALGVKHCRLP
						GITGNRIQRQATQYRDRRGTGQ
		1				NQCMRGDMQQLFADIANVAN
						TFAQIIAGRGRKFGANLLHIAD
25807	56175	В	25948	1	3738	
25808	56176	Α	25949	1518	2124	DCFCGSRHDKLPSAHQCSAKPG
						SDRFWRSHADAAYAHDGGIQP
	ļ	1				VLCASRSNGCLQKPKCHTIVSG
						RCPLQQIGS\SVFPSGDAANVAS
		1				FSCACDNAGSAQHRTLLLGKQ
						FFNCRNAFHFNRNSRQTALKSG
						ISWVTFQLILKQQPRPVIMTKM
						AIGFRKV*SQRNIIRHLLVYIFKF
						ADGGCFVVIFLCLQRFIETGINN
					<u> </u>	KMG
25809	56177	C	25950	11	747	CAMPANA AND AND AND AND AND AND AND AND AND
25810	56178	Α	25951	168	332	CAWRAVCAAW/PTGASC/CNN
						DRFVFQILRSFEHDNSEDKPGP
						GMPPHGWRTQFWGG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	I .	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25011	56170		25052	11	2706	
25811 25812	56179 56180	A A	25952 25953	1	1062	
25812	56181	A	25954	1	855	
25814	56182	C	25955	1	2412	
25815	56183	A	25956	279	713	RCDAAAGAVLHSAGLLRHCGY
23013	30103		23730	2,7		WQRGKPDRWSRRPGNYADRIC RRWFCAGGVGDRQYELCQLLA
ļ						YTVSATRRGTGYCLYRDSRGR
						TGLPVV*HLSGAGLYGRCRFAG
		1				VRWCVRHYRRTATSGIPAGDY
		ŀ				GGRVRGRNAFCHPAVAP
25816	56184	Α	25957	1	769	MDFAETELEFIKFWGDIITNKLN
23010	30104	^	23737		1,0)	EALAAQGDNVVINLASDEYFKS
						VKPKKLNAEIIKPVFLDEKNGK
						FKIISFYAKKARGLMSRFIIENR
						LTKPEQLTGFNSEGYFFDEDSSS
						NGELVFKRYEQRAECRIVTRHL
						SFSGDVCLDHDRRDDHEELSY
						GGAAVAAVLVIEPNKLAFDGS
						GYLAWEGLICMQEIGKC/YRRT
						SGDCA*VAGRAQTG*VSGWAL
		l				SRGKNQPGVGCEAGARRHARE
						ARIGDERHRLAPROML
25817	56185	В	25958	1	2472	
25818	56186	Ā	25959	4300	4492	DCGGVRSQRPDQR*DWNR*GA
						GGESDS*SLATGGQAAEGVAV
						WRYSARWR*PLFAGRCARAGG
						D
25819	56187	В	25960	49	3000	
25820	56188	Α	25961	266	812	LLRWRPGALALGHKLVAGEIR
						QQRMDAWRAACLQNPQGILCC
						ARGGQRSHIVQSWLHAAGIDY
						PLVEGGYKALRQTAIQATIELA
		1				QKPIVLIGGCTGSGKTLLVQQQ
						ANG/VDLEGLARHRGSAFGRTE
						DELGKLAQTSTSSPAHWRETSK
						CGGFLCPIFLTNMRTPLPVCAT
25021	5(100	 	25062	761	974	VLPYLAIATV
25821	56189	A	25962	761	874	MAAVCCIEACTVIEACCICTIC
25822	56190	A	25963	2	304	MAAYGSGIFAQTYIEAFGISTIQ
						PASCSPVNQYITAAFVLFGNVA HALLIALQRRNRRHLQRRKGA
						VIVIAFNTSQGA/VPAFCCQP*S
						RYASPPCCSFSTA
25823	56191	A	25964	109	777	KTASFFCCSF3TA
25823	56192	A	25965	5	187	ITHKKPPPSWRSALLPAAGVFS
23027	30172	 		ľ		RRYMPALCE*HWAS*RMRAEA
						TETFSDCTMPIIGMMMS
25825	56193	A	25966	1	711	
25826	56194	A	25967	757	1026	-

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 ' '
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25828	56196	Α	25969	1	1065	MKVLEKRNNASSEPEEKNEIRA
						AREHAGFTLQELVCLGASGRK
						VQGEAANAEEEVAAGELDNIA
						KITNEGSYTEQQIFNEGYMWLS
						TLMKIKENKISCSWATQSHFQG
						SIDAGELEHLTPERVWKETESA
ļ						LTTRNPQVFFQVLRDCGALRVL
						FPEIDALFGVPAPAKWHPEIDT
						GIHTLMTLSMAAMLSPQVDVR
			:	1		FATLCHDLGKGLTPPELWPRHH
						GHGPAGVKLVEQLCQRLRVPN
						EIRDLARLVAEFHDLIHTFPML
						NPKTIVKLFDSIDAWRKPQRVE
						QLALTSEADVRGRTGFESADYP
						QGRWLREAWEVAQSVPTKAV
						VEAGFKGVEIREELTRRIAAG
						LRG*VSRSHPHLPNAEPKNH/AS
						NYLIPSMPGVNRSVSSNWR*PA
						RLTCAAE/HGFESADYPQGRWL
						REAWEVAQSVPTKAVVEAGFK
ļ			·			GVEIREELTRRRIAAGLRGMPT
25829	56197	A	25970	2373	2597	SADDKADSVA*YLRLASA*TNS
						CRPYRRPSAPVCRRAPAI*G*TP
						QNYGPPSSCPAGGFWRYPVDPP
						RFSGFVH
25830	56198	Α	25971	2616	2777	TSVRKRFSTLSGKPVAKAGRKR
						PPPCR*TPTANGHPKPCRPPVW
					:	CHPAPVARP
25831	56199	В	25972	1	1449	
25832	56200	Α	25973	1076	1477	CAVRRSITWCKLSSKTPVRPLR
						PSTYRDCLTVSIALTPPASEKVK
						RRPVPAVKSAMKVPPGAAQSR
		1				KRVWLVSPFPTCEFLASNSRGK
		1				KISVSLSALSPRWTS*PKARWA
						ARRLTTNLVVRH*TATSVLMK
						KK
25833	56201	Α	25974	1	3552	
25834	56202	Α	25975	196	493	KRCRFADRPGDPGRQGTPAATT
	1					FGGMSFVTILPAPIPELSPMVTP
	1					GLITT*LPIQTLLPIVIGKEYISPF
	}					FLNSWCTGCPVTERVTLGAIST
	<u> </u>					LSPIVTV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25835	56203	I _A	25976	1	3917	MTDKHQQHRYVAAPGPFVGD
			<u> </u>			KCQVAGDPLMLRRALSNLLSN
						ALRYTPTGETIVVRCQTVDHLV
						QVIVENPGTPIAPEHLPRLFDRF
		1				YRVDPSRQRKGEGSGIGLAIVK
						SIVVAHKGTVAVTSDARGTSGL
						DDFFIDVVYWVRRIKRKLSVYR
						RIPRMSYRELYKPDEQPLAIMV
						PAWNERASSAIWRAGGTTLDT
						KFLFVGLPGPILGSCKVPFEFLE
						VACPPSGGPVGPPWRRPCRGA
						AAAICASPAAIILAPT
25836	56204	A	25977	3	292	GDMQRLNSLFC/SVAYSVLEQP
23030	30201	` `	23777		2,2	DEDEGRDSINIVNPSGRPIIVSSG
						ATPAVREYAAHGFHLYGLROR
						VGEVRVDLQREQIYASKPCGST
						IGISLVVR
25837	56205	A	25978	2	181	SARRLR/RSKHRFRILNRRYSHA
2000.		1.		_	1.01	HS\MLGDMQRLNSLFC/SVAYS
İ						VLEQPDEDEGRDSINIVN
25838	56206	A	25979	1	2781	V EEQT EEEE CONSTITUTION
25839	56207	Α	25980	1	2554	
25840	56208	Α	25981	635	741	
25841	56209	Α	25982	91	501	LWIRLAKRSGKAIKERRDFIFRL
						CTHRVRQGTALLRLFQHSGGN
		ŀ				RLMKLDNQHLRLRISNHRAFHF
]				RHRTTVKQSQQTFRYPWLYRV
		1				RHWGQKPKTDNFHGLLARIFG
		l				NSRSKTWGENR*EHSVLEQPDE
						DEGRDEY
25842	56210	Α	25983	2153	3062	APVMPPERLPVIAVRQYPTANC
1						CSDCLAAASRFFLASVAAPEQT
						DADTRRSRVAVGRRKTRSNSA
						NQPRPRRDWRELQLMDRLARV
			<u> </u>			AQARASFAARETLRIGDAMEQ
		ĺ				MMEGLNKVMHGEPRQEKELR
						KLADDINVLYTAIKLYLARMPK
		l				EELAEEESRRWAEIIEMSLNLEQ
						ASDIVERMG\GELVDKSWAAR
						RAFSLDGLKELDALYEQLLSNL
						KLAMSVFFSGDVTSARRLRRSK
						HRFRILNRRYSHAHVDRLHQQ
						NVQSIETSSLHLGLLGDMQRLN
						SLFCSVAYSVLEQPDEDEGRDE
25843	56211	Α	25984	135	308	KLSLRLGIFPRPSLFPRPTPSPST
						PAPAS*PAPS*VSPSPRTATPSCP
						TSYSSFPCH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	:	09/540,217	, ,	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25844	56212	Α	25985	3	663	ARGKKPPRRGKEDFWDLSIATR
						VSVNPIKFPVEA*PFLKP\LQQV
						S\GPLGGRLPLPIPGNLLLQVAD
						GTLSLTGTDLEMEMVARVALV
						QPHEPGATTVPARKFFDICRGL
						PEGAEIAVQLEGERMLVRSGRS
						RFSLSTLPAADFPNLDDWQSEV
						EFTLPQATMKRLIEATQFSMAH
						QDVRYYLNGMLFETEGEELRT
						VATDGHRLAVCSMPIGQSLPSH
25845	56213	A	25986	1	1077	
25846	56214	Α	25987	1	1323	MHSGTFNPQDFGWQGLTLTPA
		İ				AAIHIRELEILDEYGIPNAWQGE
						TNFWGSTAVSIDRLAAYKDVD
						VLCFDHDNSKDMDALMATPL
						WQAMPFVRAGRFQRVPAVWF
						YGATLSAMHFVRVLDNAIGVIV
						ATALTWMNFSQALPRSQWAQ
						AAWSPDIDVIEQMIFHYSLLPRL
						AISLLVGAGLGLDVRYYLNGM
						LFETEGEELRTVATDGHRLAVC
						SMPIGQSLPSHSVIVPRKGVIEL
						MRMLDGGDNPLRVQIGSNNIR
						AHVGDFIF\TSKLVDGRFPDYRR
						VLPKNPDKHLEAGCDLLKQAF
						ARAAILSNEKFRGVRLYVSENQ
						LKITANNPEQEEAEEILDVTYSG
						AEMEIGFNVSY\VLDVLNALKC
						ENVRMMLTDSVSSVQIEDAAS
		İ				QS\AAYVVMPMRLEGSPPLGNP
						EFQKISSPCALYIALCSKARHHT
						GRYLGLCGL
25847	56215	Α	25988	211	475	GAIGVFTAGKLTRASVYHQAR
						RFMQRETVWLVEDEQGIADTL
						VYMLQQEGFAVEVFERGLPVL
						DKARKQVPDVMILDVGLPDISG
						FELCRQLLALHPALPVLFLTAR
		Ì				SEEVDRLLGLEIGADDYVAKPF
						SPREVCARVRTLLRRVKKFSTP
						SPVIRIGHFELNEPAAQISWFDT
						PLALTRYEFIVLKT\LLKSPGRV
						WSRQQLMDSVWEDAQDTTSH
						VVAFATILQGISRDLRRRILPYCI
						RCRRERCTGTALWQ

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	<u> </u>	<u> </u>				
25848	56216	Α	25989	1661	2499	QMAHDEQWLTP\R\LQTAATLC
		ŀ				NQNAPRDGSPLWLGVDLGTCD
						VVSMVVDRDGQPVAVCLDWA
						DVVRDGIVWDFFGAVTIVRRHF
						DTLEQQFGRRFSHAATSFPPGT
	ļ	l				DPRISINVLESAGLEVSHVLDEP
	ł	i				TAVADLLQLDNAGVVDIGGGT
	1	l				TGIAIVKKGKVTYSADEATGGH
		l				HIFLTFAGNRRISLEEAEQYKRG
		l				HGEEIWPAVKPVYEKMADIVA
					:	RHIEGQGITDLWLAGGSCMQP
						GVAELFRKQFPALQVHLPQHSL
						FMTPLAIASSGREKAEGLYAK
25849	56217	Α	25990	915	1518	PSGHAPDLNGAPATAHQPGGR
						YPAPRSVRRATDNGYPAGDPLE
						WRSTAQVFVSSRRRVSLPNDRK
						TVLTRCWSAAH*FALPAPLCRD
	İ					RTGERKNRRFHTPTRCGRGFCP
						APAASRRCAGRRRGDASGRAY
						RRPAPSHCPVQ*RGHRKSTTTSS
						TSRRRPGQSARPYSGPEYRE*Y
	1					CSAQRFFHNDGSRPPVVRQAGR
						AQFP
25850	56218	Α	25991	1	1254	
25851	56219	A	25992	675	835	VRWCWIFPSQ*TLFQKRAGHA
				}		APSADVLHESCQHFLNLLNRLG
						RHYDFGEGEP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25852	56220	Α	25993	1	2255	MRESPTRDDSKNTVNTTIRIRIH
						NSYTEQLQIVTNSYCTKQWHV
						HHTAKPYAHIQRPATPDAQLTK
						MQEEEEVEPEPEMEAEVEPEPN
		l		<u> </u>		PEEAETESESMNQRDVRGTVRI
			•			VLKTFNNCRDTVFVAFEVNNT
						VSLLVATTDMTSGDTAIVVTTT
		1				GFAVFFQQRSKSAANEVQILAF
						FQGDVSFFPVATTTDTLSVTFN
						FPFNYQGVNDFDFDFKQFLHSS
						FDFCFGRVFSNFEYLHGARAHA
						CIPPSACIGEVLETSSFDIRVSVY
		l				RSSCVCVRLRESSTGDRYFGGL
				:		SLQMMRKATGLTDITSDTPGNT
				ŀ		DDQRWHPEHGLTQRMQKEQE
		İ		ļ		LVKEPEGAARDGDGEAEAWGE
				†		VKAKSSGCVGTRASSHLFRCNV
						CELHFKESSELLQHPCTPSGERP
						FRCGECQKAFKRPSG\LRQHER
		l				THSAERPFKCDLCPMGFKQQY
						ALMRHRRTHKTEEPFKCGLCE
		1				KGFGQPSHLLYHQHVHTLETLF
		l				KCPVCQKGFDQSAELLRHKCLP
		1				GAAERPFKCPVCNKAYKRASA
						LQKHQLAHCAAAEKPLRCTLC
						ERRFFSSSEFVQHRCDPAREKPL
						KCPDCEKRFKYASDLQRHRRV
		ł				HTGEKPYKCPNCDKAFKQREH
		Ì				LNKHQGVHAREQQFKCVWCG
		İ		İ		ERFLDVALLQEHSAQHSAAAA
		ł		į		AAEGAYQQTLHHPRRQQQQY
						WQLLCARNNVKLCMHILFSPH
25853	56221	Α	25994	98	596	YSGIIMAQSKLYPVVMAGGSGS
						RLWPLSRVLYPKQFLCLKGDLT
		İ				MLQTTICRLNGVECESPVVICN
1		1				EQPRFIVAHQMRQLNELSENIIL
						EPAGRTTAPTIAPAALAANRHS
						PESAPLMLVLATDHVIANEDAV
						RAPVK\NAIPYPKKGKLVTFGIG
		1				LDGRVPRYNM
25854	56222	A	25995	3	554	
25855	56223	Α	25996	1	222	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	· · ·
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25856	56224	A	25997	<u> </u>	1043	MKVKVLSLLVPALLVAGAANA
23630	30224	^	23991	l'	1043	AEVYNKDGNKLDLYGKVDGL
						HYFSDNKDVDGDQTYMRLGFK
						1
						GETQVTDQLTGYGQWEYQIQG NSAENENNSWTRVAFASLKFQ
						DVGSFDYGRNYGVVYDVTSW
						TDVLPEFGGDTYGSDNFMQQR
						GNGFATYRDTDVFGLVPTRLHP
						ARVCCALRELKDAGADRIWYI
						ADAFRAGLSVDGVFNLTNIDR
						WFLHINRAKDKNHMIISIDAEK
						AFDKIQQPFMLKTLNKLGIDGT
						YFRIIRAIYDKPTANIILNGOKLE
		İ				AFPLKTGTRQGCPLSPLLFNIVL
						EVLARAIRQEKEIKGIQLGKEEV
						KLSLFADDMIVYLENPIVSAQN
	:					LLKLISDFSKVSGYKINVQKSQT
						FLYTNNRQTESQIMSELPFTIAS
		İ				RRIKYLGIQ*IRY*WDVFQNNK
						SYL*QTHSQYHTEWAKTGSIPF
						ENWHKTGMPSLTAPIQHSVGSS
						GQGNQAGEGNKGYSIRKRGSQI
						VPVCRRHDCLSRKPHRLSPKSP
25857	56225	Α	25998	1	3339	
25858	56226	A	25999	1440	1880	AAQHASVYLYQNRIRGAVPPV
						RFFHADQRTRRDGADGKQRHA
						TETLCRIAEKISSSREQDWLHCD
						ERQSLYEWTW*PYSPHSGRSQK
						EKRQT*TAILDAYCADCFRCGR
						GVFARRERRTKRDRPGNAGTG
						GDCPCWLRRQYECVRL
25859	56227	A	26000	1	2277	
25860	56228	Α	26001	1	1845	
25861	56229	Α	26002	56	113	QQQPNPTRDQLQLAILHNIRGIR
						KIYG*IQSATREAG*GHSPQY
25862	56230	С	26003	1	2317	
25863	56231	Α	26004	3	2168	
25864	56232	A	26005	670	2541	
25865	56233	Α	26006	98	944	YSGIIMAQSKLYPVVMAGGSGS
						RLWPLSRVLYPKQFLCLKGDLT
						MLQTTICRLNGVECESPVVICN
						EQHRFIVAEQLRQLNKLTENIIL
						EPAGRNTAPAIALAALAAKRHS
						PESDPLMLVLAADHVIADEDAF
						RAAVRNAMPYAEAGKLVTFGI
					1	VPDLPETGYGYIRRGEVSAGEQ
						DMVAFEVAQFVEKPNLETAQA
						YVASGEYYWNSGMFLFRAGRY
1						LEELKKYRPDILDACEKAMSAV
						DPDLNFIRVDEEA/FLACPEESG
	. [<u> </u>		<u> </u>		DYAAMGRTGDAVVPLSLGAH

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25866	56234	Α	26007	3	417	
25867	56235	A	26008	507	717	WPEAEYLDNADAVLAMYHD/Q GLPVLKYQGFGRGVNIT/LGLPF IRTSVDHGTALEWRTCKPMVP DSIVSCIL
25868	56236	A	26009	5	378	MAVCSSLSPRRSPAFL/HDFTDV WTHHLVFGVNP*RGLIHGTALT DHVG/ENGNYSSD*IHGTALTD HVGKTETILPTKYANIYSINTSK DRFLTKPAFAEVVLFLHPRELQ TCSFFLSQNMLVKVGHQRHWF TEVDVTLTEITQLA
25869	56237	A	26010	I	469	MQKEYAYWMDGLKTCKPDNR KNALSNFGWYPSQPLLGRSRY ATTESWVEDIATAKSNPNRLPL KFTATCALPLRLAGFQ\PRWMD NPQQLNTLAPPASYRSI*TA*CL K*KNPRPRQQSCRR*RDGNQYE TLANARQKGIEKTCGTINKAG MPITT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25870	56238	A	26011		1854	CGGTLARACDGCLNGEFAA/LI TGPVHKGVINDAGI/PFTGHTEF FEERSQAKK/VVMMLATEELRV ALATT/HLPLRDIADAITPALL/H QPKYLDNADAVLAMY/HDQGL PVLKYQGFGRGV/NITLGLPFIR TSVGHG/TALELAGRGQ/ADVG SFIRPYSAIK*L*NAGASVRWLS ERRICRAITGPVHKGVINDAGIP LPVIPSFSKSVRRKRCDDAGD RRTSRGAGNDALPLRDIADAIT PALLHNRNILITPTPCWRCTRSG SSRAKIPGLRARCDITLGLPFIRT SVGHGQPLNWRDVANRCWQF YTPLFCHQMIVKPMHNRIPRAT YRGAVKSAIGLGCCCLKASATR CAYRWRRSGRRDQSRFRYFEIA GIRSRGINFIACPTCSRRNLMLS VRLTRWSNAWRYHHSDGRFD YRLRGECPGEALVSTLGVTGG NKKAASMKMGAQRPSGQQDM IDQLEARIRAKASQRTKRVELT FSRLKNNNVEKQDTALFALYQ RRLRCVRKVRRYVMAAVSAGC GGGVCAVSAYMPLNRLPNAQE ANNEYTSFENCIKCTVCTTACR
						VTLRNNDPKLPYLVKAGLKMK RVTRSPRRYCAAASATHRFNDE RAGKVQGMPDINKLPADRETC STLTCARSRQNQTNPYPANRAA NAYQVVLATKALEKVSMKSP WQHKVSSPAAVRHLPSITQRRT

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide seguence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
25871	56239	Α	26012	1	1504	MGIVASASVQAAEIYNKDGNK
		ł				LDVYGKVKAMHYMSDNASKD
		ł				GDQSYIRFGFKGETQINDQLTG
		l	,			YGRWEAEFAAWTDMFPEFGG
		ļ				DSSAQTDNFMTKRASGLATYR
						KTTSSAYRWPELNLQYHRCTS
						AARAMLLSGRPVMQINERTAV
		İ				RRQMTVYLRVERIHEIDPLVSL
						SHSMSVIPIDVERPIAAAPCTPG
ļ						LWRTKTPLRVPSTYPLRLPGRA
						RVVCHRTFRLHLCKDWVFMFS
						GLLIILVPLIVGYLIPLRQQAAL
						KVINQLLSWMVYLILFFMGISL
						AFLDNPRPYLADSVPDRQLLAP
						VNSISRYKTIEWLNYIATELHK
						GFTPLFRPDTPEEYKPTVRAHSE
						KKLQYVNEALKDEHWICGQRF
						TIADAYLFTVLRWAYAVKLNL
	1					EGLEHIAAFMQRMAERPEVQD
						ALSAEGLKGSRHLSAGARRLIL
						GIIVTFSLILALICVTQPFNPLAQ
						FIFLMLLWGVALIVRRMPGRFS
						ALMLIVLSLTVSCRYIWWRYTS
						TLNWDDPVSLVCGLILLFAETY
						AWIVLVLGYFQVVWPLNRQPV
						PLPKDMSLWPSVDIFVPTYNED
						LNVTPDELKQVLDVAAALKAL
						RAENISTKVFNSGLGISVFRDNS
						TRTRFSYASALNLLGLAQQDLD
						EGKSQIAHGETVRETANMISFC
						ADAIGIRDDMYLGAGNAYMRE
	560.46	_	26012	2000	2001	VGAALDDGYKQASGFSEPAMR
25872 25873	56240 56241	A	26013 26014	2903 123	3024 335	
25874	56242	A	26014	2634	2787	
25875	56243	A	26015	1477	2485	NPEQLMVKTQRVVITPGEPAGI
23673	30243	Α	20010	14//	2403	GPDLVVQLAQREWPVELVVCA
;						DATLLTNRAAMLGLPLTLRPYS
		ı				PNSPAQPQTAGTLTLLPVALRA
						PVTAGQLAVENGHYVVETLAR
						ACDGCLNGEFAALITGPG\HKG
						VINDAGIPFTGHTEFFEERSQAK
						KVVMMLATEELRVGLATTHLP
						VRDIADAITPALLHEGIAILHHD
						LRTKFGIGEPRILV\GGLNPHAG
						EG\GPMGTEEIDTIIPVLNELRA
						QGMKLNGPLPADTLFQPKYLD
						NADAVRPMYHDQGLPVLKYQ
						GFGRGVNITLGLPFIRTSVDRGT
						ALELAGRGK\ADVGSFITALNL
						AIKMIVNTQ
<u> </u>	<u> </u>	Ь	L			LUMANIA IA IA

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25876	56244	В	26017	1	3432	
25877	56245	A	26018	1385	2536	KSSQSHLITGMLEHVFVKSGDT
25077	30243		20010	1303	2550	PSR/LPELMNGAQALANQINT/F
						VLNDADGAQAIQLGAN/HVWK
						LNGKPDDRMIED/YAGVMADTI
						RQHGADGL/VLLPNTRRGKLLA
						AKL\PDASRTGETH/TVEWQAP
						AVAITRTATQARQSNSVDLDK
		1				ARLVVSVGRGITDSQTLRLYYI
	İ					RLPPSLMGMRLTVFLLPICGFH
						RVGPPTLLSVLATNRNRNAKVF
						AHKNAPKFRSHTASEQWKSSIA
						HNKGVHPRVTSHTHARHSLME
1						TDIPHNIHSGHASPADITRHRAQ
		1			ļ	VFNSVPKMASPIGDRVPTLHTQ
		l				HNSPANTGKAQSLLTCGPAAES
						CCRLIPRQMVSLVFRLKARLLLI
	ļ					TTELPKGLVPVATISGTGNFGN
						TQSANPYRPSENTDLSLS
25070	56246	A	26019	926	1293	LSPNPKEPSCRTRLKKSGFFTRI
25878	30240	A	20019	920	1293	SAGAEPLMAIPARLSNSLACGA
						RDSAAKE/GFVVVPVCVAVEA
						DIAELDDEERDEFMADVGFT\V
					,	GVKEVRAWTIPVGATAPQAAG
						KIHTDFEKGFIRAQT
25879	56247	A	26020	1	1278	KIII DI EKOI IKAQI
25880	56248	A	26021	678	1526	ERNTNTCEVVPNRKRAYASHD
-		-				MELSISPVGKGASKGEVNPADD
		1				IEVINTELALADLDTCERAIHRV
İ		ŀ				QKKAKGGDKDAKAELAVLEK
					ļ	CL\PQLENAGMLRALDLSAEEK
						AAIRYLSFLTLKPTMYIANVNE
						DGFENNPYLDQVREIAAKEGSV
						VVPVCAAVEADIAELDDEERDE
					1	FMQELGLEEPGLNRVIRAGYKL
						LNLQTYFTAGVKEVRAWTIPV
				,		GATAPQAAGKIHTDFEKGFIRA
						QTISFEDFITYKGEQGAKEAGK
						MRAEGKDYIVKDGDVMNFLFN
25881	56249	Α	26022	1177	1605	
25882	56250	A	26023	1	644	MDNHIALDFLLASECGSVYVY
					1	ATTHGRNMAGARALWRATGM
						TDADFGKPIIAVVNSFTQFVPG
						HVHLRDLGKLVAEQIEAAGGV
						AKEFNTIAVDDGIAMGHGGML
						YSLPSRELIADSVEYMVNAHCA
						DAMVCISNCDKITPGMLMASL
						RLNIPVIFVSGGPMEAGKTKLS
						DQIIKLDLVDAMIQGADPKIN/D
						YQSRGRRCPNRCFRCKNQAGS
25883	56251	В	26024	1	1944	
23883	20221	lR	26024	11	1944	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25884	56252	Α	26025	1 '	1488	
25884 25885	56252	A	26025 26026	1	1488	MNKAHFVSNLGATKCCPLSFLP YRWRRVIAFVVAFYFRHLIDLL VVQMHFKLVKGNVDVVFTAQ LIDSTVQLIDSTTTVTQFTHIQT QLMIQRAGTKCRKEDFRHCVGI LEPEWYPIIVVRVVVDVSQFEV RSQSLIFAVGEGITCIYTQFITHA VVRLTVNFHVVAQLSSPDWNR PETLPVRSHVTSITQFEVVTGFQ TEAPSVGVGATVNVSTVAHGC RQVVVEVRVAQATINKDVVRE VSRRVDVGSLAVLIHLTRTVIH VTFSEACSRTDDPLSVVTVNFGI NTDHDAVNILIIVDGAVVAVEV TTEVAYPCAAVISQAMTRVGQ TSTNGVWTISSVNAFQIRTGAA RSLAPPWSLTEKVFCGTTMER MRAALEAIMPFTESSSATQAPY LPCGRECGLAVRQFFGRGEGIE TVAYLFRIIRVVRTRVLSIGGFS RTRISSTPVKIATLRQNRSDCHA AMIADTYRQVTPTDPFSRHPFS
						RVLRGLFFIIFKDTKAHNGIGMI GGVNRRADYLNLLEKVFGDRV SSPQWQATLKAIVPSYGRKLNG DVAATERELQYTSEVVGLNYD KPQAADNNRQHKVRAARSRRQ NRPDRGSQIADSTAGARCRFQT PTAGDGA*SAASTAAPI/MLNLL EKVFGDRVSSPQWQATLKAIVP SYGRKLNGDVAATERELQYTS EVVGLNYDKPQAADNNRQHK
25886	56254	Α	26027	1	2130	
25887	56255	A	26028	1144	1309	PGGSTPMAVEDPDFRSSERVLR FQPAVSVYGPEQPAV*DYAQTS YLRTRPRRSDP
25888	56256	Α	26029	1	3546	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25889	56257	A	26030	1097	3332	TRVLVLDHRLLRCLYATVCAG
						ASWGMNPVVCSQQIDPQFHTM
						LMICASEEAQRMQQASEDEFN
						RALNIAFDNRLGLCKVESARQV
						FPLTGRYARQFASHRLALVGD
						AAHTIHPLAGQGVNLGFMDAA
						ELIAELKRLHRQGKDIGQYIYL
						RRYERSRKHSAALMLADSFVR
						TTHALRRSHGGFPRLDDAAAL
						RFANRRTSCERNDDNTWSRSL
						YADLKNGTAQNIRAKFVFIGAG
						GAALKLLQESGIPEAKDYAGFP
						VGGQFLVSENPDVVNHHLAKV
İ						YGKASVGAPPMSVPHIDTRV\L
						DGKRVVLFGPFATFSTKFLKNG
						SLWDLMSSTTTSNVMPMMHV
	-					GLDNFDLVKYLVSQVMLSEED
						RFEALKEYYPQAKKEDWRLW
						QAGQRVQIIKRDAEKGGVLRL
						GTEVVSDQQGTIAALLGASPGA
						STAAPIMLNLLEKVFGDRVSSP
						QWQATLKAIVPSYGRKLNGDV
						AATERERKTLHVIFNDLRPEQC
						EAGKSPSRVQLQTWTNGGMLN
		1				APLSLRLTLVEKLASMLDPGHL
						ALTQIAQHLALLQKMDHRQHS
		1				AFPELPQQIAALYEWFSARCRW
	İ					KEKALTQRGLLVQAGDQSEQIF
						TRWRAGAYNAWSLPGRCFIVL
						EELRWGAFGDACRLGSPQAVA
						LLLGDLLEKATQHLAESINAAP
			:			TTRHYYHQWFASSTVPTGGEH
25890	56258	Α	26031	115	654	HATQTMTPWFLYLIRTADNKL
						YTGITTDVERRYQQHQSGKGA
						KALTGGSARSPLIKKALAEQLP
						GIPIAGGDDFGSVTAGLAR*RF
						HQRHGPCWSMKAESLQLCRAD
						CQRRFQPMSFGGHFRHYRLSVC
		1				WRYRFIRPEHKTAFTARFPVDN
						RLGQMVAKPFHLLNATTGNIC
						GCFNTKRP
25891	56259	A	26032	1	348	
25892	56260	Α	26033	1	906	
25893	56261	В	26034	59	910	
25894	56262	Α	26035	1	3684	
25895	56263	A	26036	1	2634	
			<u> </u>	٠	<u> </u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	•	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25896	56264	I _A	26037	1162	3478	LVRFGLLPLAYSARCFSARSED
23070	30201	``	20037	11.02	3 77 0	RPKDECETCCIKYPNGRNVLSQ
		İ				EIHQVFVLNGIQSMSGYVYNLG
į						NELASMQGLVDVVRLSPQGSS
						AREAFSHSLDLARLAEKRGYH
		ŀ				RYWLAEHHNMTGIASAATSVLI
į		İ				GYLAANTTTLHLGSGGVMLPN
						HSPLVIAEQFGTLNTLYPGRIDL
						GLGRAPGSDQRTMMALRRHM
						SGDIDNFPRDVAELVDWFDAR
						DPILEPTHNGRRLCSTVRHPRM
		ŀ				
ì						GYTPGVEPTTGPLGQGLANAV GLAIAERTLAAQFNQPDHEIVD
						HFTYVFMGDGCLMEEGWFTD
						DTAKRFEAYHWHVIHEIDGHD
						PQAVKEAILEAQSVKDKPSLIIC
						RTVIGFGSPNKAGKEEAHGAPL
		ŀ	i			GEEEVALARQKLGWHHPPFEIP
						KEIYHAWDAREKGEKAQQSW
						NEKFAAYKKAHPQLAEEFT\RR
						MSGGLPKDWEKTTQKYINELQ
	[ANPAKIATRKASQNTLNAYGP
						IMLPELLGGSADLAPSNLTIWKG
						SVSLKEDPAGNYIHYGVREFG
						MTAIANGIAHHGGFVPYTATFL
		ŀ				MFVEYARNAARMAALMKARQ
						IMVYTHDSIGLGEDGPTHQAVE
						QLASLRLTPNFSTWRPCDQVEA
						AVGWKLAVERHNGPTALILSR
		l	}			QNLAQVERTPDQVKEIARGGY
		1				VLKDSGGKPDIILIATGSEIEITL
						QAAEKLAGEGRNVRVVS\LASS
25897	56265	A	26038	1	1033	MNKSPKTRDIQEEELEVLLEEE
23697	30203	^	20036	1	1000	AAGGDEDREKEILIERIQSIKEE
						KQKGLGLAAGWTADVSSRRPG
						ASQLPPPPTHPIPQYRGWGPPD
			İ			QHLLSTLWTNVGSYKSGSFLQL
						GGWCMNLDMDLNSEESRWLF
						RRHWDTGGIKPHERKRAVAHL
						TLSEREEIRAGLSAKMSIRAIAT
		1				ALNRSPSTISREVQRNRGRRYY
						KAVDANNRANRMAKRPKPCLL
						DQNLPLRKLVLEKLEMKWSPE
						QISGWLRRTKPRQKTLRISPETI
						YKTLYFRSREALNHLNIQ\HLA
						RSHWLWPGERY\NRNGERGPV
						TLWTG/TPISDGSRTSVKRALGQ
						WGEHLSRRNAFFPLCATVV
	}		l	J		WUEDLSKKINAFFFLCATVV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25898	56266	Α	26039	797	1440	 PAMRLTAMRCISMKSRLPASR
						MQFCATVRWKYRCHREPTLSA
						QHGAFRSINPTPRNSRGNCWKK
		l				REMKWANWPIWCR*KN*RERS
						LSAICCASRK/TGQFVLPPARYH
		1				LRRSHSLRHGRRHTRKGERGTI
		1				NIVNGTPIHERSRNIDNRRSLGH
		1				WEGDLVSGGVGTVIGVVTGVII
		1				FTVINYGLTYIGVNPYWQYIIK
		1				GAIIIFAVALDSLKYARKK
25899	56267	A	26040	687	1106	GSLSLTALLPDTAEREEIRAGLS
230))	30207	` `	20010	007	1	AKMSI/RAIATALNRSPSTISRE/
						VQRNRGRRYYKAVDAN/NRAN
						RMAKRPKPCLLDQ/NLPLRKLV
		ŀ	ļ			LEKLNIQHLRRSHS/LRHGRRHT
						RKGERGTIN/IVNGTPIHERSRNI
		İ				DNSALVTVLR
25900	56268	A	26041	1	2851	MKTLIARHKAGEHIGICSVCSA
23900	30208	^	20041	1	2651	HPLVIEAALAFDRNSTRKVLIEA
1						TSNQVNQFGGYTGMTPADFRE
1						FVFTIADKVGFARERIILGGDHL
			:			GPNCWQQENADAAMEKSVEL
						AVVIDLWSRAVIGWSMSPRMT
		ŀ)
1						AQLACDALQMALWRRKRPRN VIVHTDRGGQYCSADYQAQLK
1						RHNLRGSMSAKGCCYDNACVE
						SFFHSLKVECIHGEHFISREIMR
						i
						ATVFNYIECDYNRWRRHSWINS
25901	56269	В	26042	1	1048	LLSQKRNTQGRIEDGRQH
25901	56270	A	26043	323	1290	GTSMNTVGTPLLWGGFAVVVA
23902	30270		20043	323	1290	IMLAIDLLLQGRRGAHAMTMK
1						QAAAWSLVWVTLSLLFNAAF
		l				WWYLVQTE\GRAVADPQALAF
						LTGYLIEKSLAVDNVFVWLML
1		l				FSYFSVPAALQRRVLVYGVLG
1						AIVLRTIMIFTGSWLISQFDWIL
						YIFGAFLLFTGVKMALAHEDES
						GIGDK\RWCAGYA/VHLRMTDT
1					}	
				1		IDNEHFFVRKNGLLYATPLMLV LILVELSDVIFAVDSIPAIFAVTT
]
						DPFIVLTSNLFAILGLRAMYFLL
			}			AGVTKRFSMLKYGLAGILVFIG
						IKRLIVNFNLLQTRACLGGGWA
		<u> </u>				FGETLIIKPGEYRMKTAGG
25903	56271	A	26044	28	96	
25904	56272	A	26045]1	777	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	tocation of first codon for peptide		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
25905	56273	A	26046	77	532	KPSRPEISKGKSVKPAASRRARE
23303	302.3	ļ^^	200.0	' '	332	CAVQALYSWQLSQNDIADVEY
						QFLAEQDVKDVDVLYFRELLA
	ļ 					GVATNTA\YLDGLMKPYLSRLL
						EELGQVEKAVLRIALYELSKRS
		1				DVPYKVAIKEAIELAKSFGAED
						SHKFVNGVLDKAAPVIRPNKK
25906	56274	Α	26047	3	175	
25907	56275	A	26048	85	440	YSSSSLWAARKRRQMVLSARR
						FATSDRVCRCVLVWSAGTSRA
		İ				KSKSTGWLSMASKAIGVSS*TK
						TPTARRVSSSSLPCGMAIPLPIP
						VLPIFSRVRIASKTT*GSSLSCFA
						ARSLMTS
25908	56276	Α	26049	80	706	KLIQVAVVIGGGQPLGAFLCHG
						LAAEGYRFPVVDIQSDK\AANV
						AQEINAEYGESMAYGFGADAT
						SEQSVLALSRGVDEIFGRVDLL
					:	VYSAGIAKAAFISDFQLGDFDR
		1				SLQVNLVGYFLCAREFSRLMIR
						DGIQGRIIQINSKSGKVGSKHNS
		1				GYSAAKFGGVGLTQSLALDLA
		l				EYGITVHSLMLGNLLKSPMFQS
						LLPQ*KTKLGIK
25909	56277	A	26050	741	893	TGCSGKRSLQQRREERRRNPVQ
						KIRCTV*RVRNGRIRYCSPRAGF
						YSRYR
25910	56278	A	26051	269	987	MD A V CD D CD DDWG WD DD C
25911	56279	Α	26052	1	1268	MDAKCDRSRFPWLWPTKRPRG
						CFATFVPIGIPYDQTKTHLHTLS
						LVAKRLADKTICRRLRAAQSDE
						ELYQIITDTEGTPDEARCEEKQY
						MVLMIVSGRSGSGKSVALRAL
					•	EDMGFYCVDNLPVVLLPDLAR
						TLADREISAAVSIDVRNMPESPE
						IFEQAMSNLPDAFSPQLLFLDA
						DRNTLIRRYSDTRRLHPLSSKN
						LSLESAIDKESDLLEPLHGFPDY
						DTVGFSCKRRIDYVCRIKHSRRI
						RQVVLLNFAKSGAFSTTRGTDD
		1				KTRRSLLVTL/VAYFLCAR/EFS
		1				RM/MIRDGIHVRIIQINSKSGKV
						GSKHNSGYSAAKFGGVGLTQS
		1				LALDLAEYGITVHSLMLGNLLK
						SPMFQSLLPQYATKLGIKPDQV
						EQYYIDKVPFKRGCDYQDVLN
25012	56000	 	26052	1	160	MLLFYASPKA\SY\CTGQSINVT
25912	56280	Α	26053	1 .	168	LKVFILVIFTLSRE*EHAMFGFL
1						FLC*FAENDGFQLHPCPCKGHE
<u> </u>	56281	A	26054	2	253	LILFYGCIVFH
25913						

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25914	56282	В	26055	319	3780	
25915	56283	Α	26056	513	762	
25916	56284	A	26057	375	550	GVGLCGRSSIRRNM*CK**G*/L SPKIALMACGRRHTRSLFMTSK DLSRMMRLQKEIRL
25917	56285	A	26058	662	1033	EPKRASGCMWPVGAHCHHPLL LLACQQKLGPPSHQASSRS*NIQ TLDRLCMSSWIPQEHSSSQSWH RSCFLNVTRPWSFYSPVGKNR WMGPFWSFSLDALLLFFTLHSY VAWPVEMMALNLLG
25918	56286	В	26059	1	2034	
25919	56287	A	26060	369	515	HAKPPGGLGHCSLCPSHYSCIH G*KGPRYVSVSCSRGWKPEAT KAFMW
25920	56288	Α	26061	352	2631	
25921	56289	В	26062	10	351	
25922	56290	A	26063	321	537	EPLDTGRGTSHTRACHGVGYW GRDSVRRYT*CK*RINGYSTPT WHMHEYVTNLHIVHMYPRTYS IIIKKMSD
25923	56291	A	26064	788	912	VHTHLSERTRKCLPPPAPS*WV TCHHGDCCLPTRLRAPQGP
25924	56292	Α	26065	1348	1799	
25924 25925 25926	56294	A	26065 26066	543	741	ATYPSAHYPDHNRGEFLWPSL GLGKELDTFWSSLRVFGSPVSA VSRELVRKFSLGGTVDSTGSLP VCGDISPVSDVA/ACTREPLRFR LQVGDRYITLMDLPGVGESGA RDTEYAALYREQLPRLDLVLW LIKADDRALTVDEHFYHQVIGE VYRHKVLFVISQSDKAEPTSGG GQLSTAQKQNISRKICLLHELFQ PVHPVCAVSVRLQWGLKVMA ERMIKCLPREATSPVVSQLHPSF RTTVVREQARSDFGETVGAVL DSISAFPLIPAPVRAVIQAVRTT VVSFAASVTLLLLPLRSTPVEKS PTCILPASETFPLLVVMLNPVAS LLTLTSPVLDTSCPTHSLSAAQP NLIQLSGNNNISFVVNRAVVTN VLNTMRTACNSQATIQIYNADI ARDFGTRGIFSINSGFSTVYSRRI GHQSPEVTGEAMALTLSSQPSC IDDGLVAVNLDTDGLRIGDARQ KAITAGYPVVDTDDKCRIRRLR RIRHEQCADENSYQAYNYCRP VGRIRHLCRIRQSMPDATLSRLI

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
25927	56295	A	26068	3	373	VYTGLKPRTIDAQYTPINVV/W KFLSLTHHAYSPLLVVINKAKF DGLSPEFQQALVSSAQEAGNY QRKLVAEDQQKIIDGMKEAGV EVITDLDRKAFSDALGNQVRD MFVKDVPQGADLLKAVDE
25928	56296	A	26069	I	1107	MGAFTGKTVLILGGSRGIGAAI VRRFVTDGANVRFTYAGSKDA AKRLAQETGATAVFTDSADRD AVIDVVRKSGA*ILILKRRSISSA FNSRASPKTPIPAFTTRISNAPLL RTTSMTASLSALSVNTAVAPVS CARRLAASFDPA
25929	56297	Α	26070	39	254	
25930	56298	Α	26071	1432	1845	
25931	56299	A	26072	674	1614	AVIVVAILVSKSTGAWVAFSIR MPSITFSHNSVVAGVASARKDS SPSYGDPLESAIPKLNVKRSRSG LSKSLLTLTSCSTIWINWITGQT PLKPCSVTGSVVPKAWRSPSTL TTMTTR*PFTLPA/LDTFISRPAG CTYANECHAPESATIRNDVAAR PCNCAQCAALDCVSVFVTIRHS CSSVYSAPGILVAALQAGHKPV ALVGGATGLIGDPSFKAAERKL NTEETVQEWVDKIRKQVAPLP DRKNISPADSRLSPDGRYRGYV HPAPPIPDSHLIRHHRCVIYQWR QCLAQFRLNGWAASRSQSICRI GCD MSSSCIEEVSVPDDNWYRIANE
25932	56300	A	26073		3476	MSSSCIEEVSVPDDNWYRIANE LLSRAGIAINGSAPADIRVKNPD FFKRVLQEGSLGLGESYMDGW WECDRLDMFFSKVLRAGLENQ LPHHFKDTLRIAGARLFNLQSK KRAWIVGKEHYDLGNDLFSRM LDPFMQYSCAYWKDADNLESA QQAKLKMICEKLQLKPGMRVL DIGCGWGGLAHYMASNYDVS VVGVTISAEQQKMAQERCEGL DVTILLQDYRDLNDQFDRIVSV GMFEHVGPKNYDTYFAVVDR
25933	56301	Α	26074	200	1100	
25934	56302	A	26075	1	2061	
25935	56303	A	26076	1	3294	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
-				sequence		
25936	56304	Α	26077	1	2064	MNYSLKQLKVFVTVAQEKSFS
						RAGERIGLSQSAVSHSVKELEN
						HTGVRLLDRTTREVVLTDAGQ
						QLAFRLERLLDELNNTLRDTGR
						YGATTERSARYSASLPAPTSGR
						SEPAALVQNSSSQPSSLKENIM
						KLRSVTYALFIAGLAAFSTFFLA
						\AQSIRCGYETSQADSQHSGAK
						KFNDLLQERPKGELKLKLFPDS
		1				TLGNAQAMISGVRGGTID\MEM
						SGSNYFAGLSPVMNLLDVPFLF
						RDTAHAHKTLDGKVGDDLKAS
						LEGKGLKVLAYWENGWRDVT
						NSRAPVKTPADLKGLKIRTNNS
						PMNIAAFKVFGA\NPI\PMPFAE
		l	ļ			VYTGLETRTIDAQEHPINVVWS
						AKFFEVQKFLSLTHHAYSPLLV
						VINKAKFDGLSPEFQQALVSSA
						QEAGNYQRKLVAEDQQKIIDG
						MKEAGVEVITDLDRKAFSDAL
		ł	,			GNQVRDMFVKDVPQGADLLK
						AVDEPGWAERDIWAFAPAFFY
						PLFISDFNRVRLEFVGHYQDVC
ł						EKPASTTLWLDVGRSSGLDLTY
						QTLNVKNDLSHFPVPFFDPSDN
						RTNTLPMVFAGAPDVGLQQAS
						AIVASWFGSRSGWRGQNFPVL
		ļ				YNQLPDRNAIVFATNDKRPDFL
		1	}			RDHPAVKAPVIEMINHPQNPYV
1		1				KLLVVFGRDDKDLLQAAKGIA
1		1				QGNILFRGESVVVNEVKPLLPA
						GSQPVWSSSASESRCCFYA
25937	56305	Α	26078	1	480	LGNTKTVKGWLAQLPAKYHQ
					ł	RATCMFDRHGLLALLAGRFLA
						FVRTLLPTMAGISGLPNRRFQFF
İ						KAKLR\WLS*PTRIAAALAFMP/
						FMDDHHPRLMQTVILQILQRRG
						TGDRLKIMVERRYAHVGFCRQ
						LLDAQVFGVFILNPFQHAANQT
						EVSLATDQRQ
25938	56306	Α	26079	1	2184	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25939	56307	A	26080	1429	2152	NLFREVVFPTHCPVCGSDVERV
			İ			EGEAVARCTGALICGAQRKESL
						KHFVSRRAMDVDGMGDKIIDQ
						LVEKEYVHTPADLFKLTAGKLT
						GLERMGPKSAQNVVNALEKAK
						ETTFARFLYALGIREVGEATAA
				-		GLAAYFGTLEALEAASIEELQK
		ŀ				VPDVGIVVASHVHNF\LPKKAT
		l				AMSSASCWRKVFTGLRRSLSTR
						KRLTARLLVKPWCLRAASGTP
•		l				LKRASLLMFSGCFSHTSGLAMY
25940	56308	A	26081	1	2448	
25941	56309	A	26082	125	741	AHLNRTRWKVCWIR*RKISPW
		`				AKRNTKTVKG\WLAQLPAKYH
						QRATCMF\DRHGLLALLAGRFL
						AFVRTLLPTMAGISGLPNRRFQ
		ĺ				FFNWLSGLLWVSVVTSFGYAL
		İ				SMIPFVKRHEDQCRNGVFGKR
				:		KGGIAAALILTRLRRRFVSSWK
						RFRELRRKSSQADDMNVFGILE
						DFTHQFIGGAFLRADISGAAKM
		l				SCLAKAQASSSMH
25942	56310	A	26083	1066	1359	
25943	56311	Α	26084	256	952	RLKRCSTGKSMAVIQDIIAALW
			ŀ			QHDFAALADPHIVSVVYFVMF
ĺ						ATLFLENGLLPASFLPGDTLLIL
						AGALIAQGVMNFSGLRIAVLDP
						RNQVRGGGLSIFKGRWLGNPK
		1				TVKG\WLAQLPAKYHQRATCM
	ļ	1				FDRHGLLALLAGRFLAFVRTLL
		1				PTMAGISGLPNRRFQFFNWLSG
	İ	1				LLWVSVVTSFGYALSMIPFVKR
1		1				HEDQVMTFLMILPIALLTAGLL
		1				GTLFVVIKKKYCNA
25944	56312	В	26085	1	1290	
				<u> </u>	<u></u>	<u> </u>

SEQ ID			SEQ ID NO:	1	•	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
25945	56313	I A	26086	1	1839	MRCQNA WWIMQHSSGSERLSS
23943	130313	A	20080	1	1039	` `
						VLLHFRRTNLADSNDRIQYHSL
						RFFDEDGAQWLVILGDVLNHG
						PRNALPDGYAPRKVAERLNEV
						AHKVIAVRGNLRQRVDQMLLI
						SRAINRLVASNWNRRSAQMRC
						PISGKNLAWIPVQLPVYWQNN
						CRRLLMRSHRKHGSGTNLPGE
						LMAIKLIAIDMDGTLLLPDHTIS
						PAVKNAIAAARARGVNVVLTT
						GRPYAGVHNYLKELHMEQPGD
						YCITYNGALVQKAAD/GYDDY/
						RF/LEKLSREVGSHFHALDRTT\
						LYTAN\RDISYY\TVHEFLRCHQ
İ						FPLVVCDGGRMTPRAMFLFSSA
						HILMPLNYHNSHINTLFPVAGT
						LMVEPTESESKVELDRFIDAML
						AIRAEIDQVKAGVWPLEDNPLV
						NAPHIQSELVAEWAHPYSREVA
						VFPAALVTGGSRGIGRATALLL
						AQEGYTVAVNYQQNLHAAQE
		İ				VMNLITQAGGKAFVLQADISDE
						NQVVAMFTAIDQHDEPLAALV
		1				NNAGILFTQCTVENLTAERINR
<u> </u>		1				VLSTNVTGYFLCCREAVKRMA
						LKNGGSGGAIVNVSSVASRLGS
		1				PGEYVDYAASKGAIDTLTTGLS
		1				LEVAAQGIRVNCVRPGFIYTEM
25046	56214	ļ_	26097	12	1275	HASGGEAWTRRSR
25946	56314	Α	26087	2	1275	VNFSPKSSQIHHALRTVAGRFA
						VKSIDYFWHDSCNASKRFHIWE
						SIMLELLFVIGYLVMLMVTGVS
						LLGIIAALVVATAIMFLGGMLA
		1				LMIKLLPWLLLAISVVWVIKAI
						KAPKVPKYQRYDRWQLAVRRS
						PLLPDHTISPAVKNAIAAARAR
						GVNVVLTTGRPYAGVHNYLKE
						LHMEQPGDYCITYNGAL/VYRK AAGWLAPLAAKLLKLMDELI
						· ·
						GFLEKLSREVGSHFHALDRTTL
		i				YTANRDISYYTVHESFVATIPLV
					1	FCEAEKMDPNTQFLKVMMIDE
					1	PAILDQAIARIPQEVKEKYPVLK
						APNLQQPHGSSGSSFHREIFFSS
					İ	RSEDKRRGGLVVSGISLLVVSA
				1		GPWVRNANEPLGQLKTWFSLS
						HTLTLTYTRAPACRLTSTQTEP
						RARPCLSPLSSEPRDRSLTPRLG
0.50.4=	56017	<u> </u>	26000	1	1.500	QRFIYLAVQ
25947	56315	В	26088	1	1509	
25948	56316	Α	26089	1	1323	_

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25949	56317	В	26090	1	5073	
25950	56318	Α	26091	249	531	CWAMRMDSKVGVITPIRMWSA
	1	ŀ				SGPFAMGTLVPTFAAVTDIPTK
						SVTTCLRQICERWQAPSTR*SRF
						ASLSQPAPRFTPSPITLALVLLP
						ARRC
25951	56319	Α	26092	3	1038	LVLFGVAETYPVANDFAAGAG
						AYVRGRYAAGVRAWPTIITRK
						TMKTLVVALGGNALLQRGEAL
ŀ						TAENQYRNIASAVPALARLARS
						YRLAIVHGNGPQVGLLALQNL
						AWKEVEPYPLDVLVAESQGMI
						GYMLAQSLSAQPQMPPVTTVL
						TRIEVSPDDPAFLQPEKFIGPVY
						QPEEQEALEAAYGWQMKRDG
						KYLRRVVASPQPRKILDSEAIEL
		1				LLKEGHVVICSGGGGVPVT\ND
						GAGSESVIHKDLAAALLAEQIN
						ADGLVILTDADAVYENWGTPQ
						QRAIRHATPDELAPFAKADGSM
						GPNVTAVSGYVRSPVQMHHRQ
						FQPVIIHIATNQIDHHRRIIHHWL
25952	56320	Α	26093	I	3639	MKRLIVGISGASGAIYGVRLLQ
						VLRDVTDIETHLVMSQAARQT
						LSLETDFSLREVQALADVTHDA
						RDIAASISSGSFQTLGMVILPCSI
						KTLSGIVHSYTDGLLTRAADVV
						LKERRPLVLCVRETPLHLGHLR
						LMTQAAEIGGNTGEIDEEELES
						LLYAIAKGNVFNFQTILHLPVA
						VQNDTIDFYQMFARIWSSHPQ
						WLTLYLAQHRAVIIPDDAKLHR
						NLLRWYSAGRLDIPELLDYAQS
						WRETEPDNEDAPY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25953	56321	A	26094	1	1260	AWARQDRRLSVKQWGKPPVD
	20321	1	2009 1	ļ. •	1200	ETIPAKEQVRILRKLFDRYYGE
[VAGRGDGFCSPHVARYVGRVA
						G*RPCRLGPGHPTNRTPFVAPL
1						LEALDIAKYFSVVIGGDDVQNK
						KPHPDPLLLVAERMGIAPQQML
						FVGDSRNDIQAAKAAGCPSVG
						LTYGYNYGEAIDLSQPDVIYQSI
						NDLLPALGLPIAKIRNRKMTKPI
						VFSGAQPSGELTIGNYMGCTEA
						DRQFAFNFNNFVLIQLLTNGGP
						DRLGTTTPAGYTDLLVNYTYRI
						AFEGGGQDFGLAAAIATLIFL
						LVAIMFPLLMVVAISLRQGNFA
						TGSLIPEQISWDHWKLALGFSV
						EQADGRITPPPFPVLLWLWNSV
						KVAGISAIGIVALSTTCAYAFAR
						MRFPGKATLLKGMLIFQMFPA
						VLSLVALYALFAPRVRAALCY
25954	56322	A	26095	l	3240	
25955	56323	Α	26096	494	792	ATRLRHGVTSVGGDGGLQLTV
						MDKLDVILLMSV/NPGFGGQSFI
						PQTLDKL/REVRRRIDESGFDIR
						L/EVDGGVKVNNIGEIAA/AGA
						DMFVAGSAIFDQPDAR
25956	56324	Α	26097	1624	2390	RTSERRWHAKALLKLFSAHHA
						GFVCCKEEEERLATGKVGWLP
						VETLLAQQHIEGHYQVDPSLFK
						PNADCPVRVSGMSMKDIGIMD
						GDLLAVHKTQDVRNDPLLVPL
						FTLIREGKLAANWPLEQDELLT
						RLQKSCDMTQVSADYNALFIG
						DECAVPPYRSAWVEGATEAEV
						RAFLSERGMPLADTPADHIGTL
						LLAASWLEDQSTEDESEALETL
						FKRGA/LPKSLCLWGKAPAPHG
						SPVGGFSKGPKTWAPHEGAS
25957	56325	В	26098	1	534	
25958	56326	A	26099	67	399	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25959	56327	A	26100	8	1016	PESRRKAVMTQPVLDIQQLHLS FPGFNGDVHALNNVSLQINRGE IVGLVGESGSGKSVTAMLIMRL LPTGSYCVHRGQISLLGEDVLN AREKQLRQWRG/ARVAMIFQEP MTALNPTRRIALQMMDVIRHH QPISRREARAKAIDLLEEMQIPD AVEVMSRYPFELSGGMRQRVM IALAFSCEPQLIIADEPTTALDVT VQLQVLRLLKHKARASGTAVL FISHDMAVVSQLCDSVYVMYA GSVIESGVTADVIHHPRHPYTIG LLQCAPEHGVPRQLLPAIPGTV PNLTHLPDGCAFRDRCYAAGA QCENVPALTACGDNNQR\GAC WYPQQEVISV
25960	56328	A	26101	3	102	ISRLNTQQHFG*RTRQISPHSPQ GCSRCPLHS
25961	56329	A	26102	290	490	SSLNSCMAVWDKTRLIRQ*LV MWSY*SPSPCR*PAGYRHMKL RSTSLVLSTPSRLFLAVIPPVVVI
25962	56330	Α	26103	1	3255	
25963	56331	A	26104	119	634	
25964	56332	A	26105	1640	1770	HLDSNRRVHDSTCNWCCNLS* QRLLWSG*WSELCCSVPRQKR A

SEQ ID	1 .	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ļ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
25965	56333	A	26106	281	2108	RESLKQKVGGRPQLSPFPPPDS
20,00		`	20100			WALDPGGGRARETPRGVSNAT
						KEPISVAQHGN/GGDKEIRRELK
		1				YYAHLGRRLKDWSGIPRYHGT
	,	1				VETDCGTGYVYDVIADFDGKP
						SITLTEFAEQCRYEEDIAQLRQL
	1	-				LKQLKRYLQDNRIVTMSLKPQ
						NILCHRISESEVIPVVCDNIGEST
						LIPLATWSKWCCLRKQERLWK
		Ì				RFIAQPALAIALQKDLQPQATE
						ASGAENMASIASGNIARTDPDM
						LSIPLSKKICCFGDRIIEIPETIDQ
						PVRFWVLTGPVVDMGIYACYL
		İ				SSPOWGWYAERDAEIENETVR
		İ				REVEELRQASETDLQPGTIEYE
						RHRLTRAQADAQELKNARDSA
						EVVETAFCTFVLSRIAGEIASIL
						DGIPLSVQRRFPELENRHVDFL
						KRDIKAMNKAAALDELIPGLL
						SEYNRADRQYAGGSRVAARFA
						ASPRVAVLSVITGLIPRHVCAPY
			<u> </u>			LLCGDPCCLSTPEDMTPRFRRY
						AVRSGNRSTPGLDHRCDACQR
						SAYELADLRCRPPPSTLFLLPEA
						DKNDFSDALPAVADAMYAAPP
						RRKAAPSKPEKAVVSPLRSVRK
				1		ARHGSVCSLRPDQYAVRNLPA
					ĺ	YRSCSSPDLQFMVWCALA
25966	56334	A	26107	1	2880	1 RSCSSPDLQFMV WCALA
25967	56335	В	26107	61	5101	
25968	56336	A	26109	80	561	IRPLPPRFKTESRSLPGPCLQPGT
25700	30330	``	20107		301	FLWSRNRRVLGFPSMNGEDMG
		l			}	LLFLCSEWERSSEGWLCNREGG
		1				SGHSIEPHCCTFLHLTHRSLAFS
				1	1	LLAGVSCTCASSCKCKE\CKCTS
						CKKSECGAISRNLGLWLKVGR
		İ				EPKAVPEVRASGEPAFLCPCRL
		ł				SLPV
25969	56337	A	26110	1	277	· · · · · · · · · · · · · · · · · · ·
25970	56338	Α	26111	902	1006	
25971	56339	Α	26112	2	1265	
25972	56340	Α	26113	409	525	
25973	56341	Α	26114	91	431	NHKPGNIDVARRIQRGFAGDQI
				1		GHLRPVERQCSPDKRRFIAADG
				i		REIRGKQRAGHIFQLLSRCLLQI
						LNHCQRRAAHFRFQLSNQRHQ
						QLLPV/HYHAAEREYPAGACLV
	<u></u>	L		<u></u>		RWLL
25974	56342	Α	26115	3	71	
25975	56343	A	26116	li .	1212	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	3	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
25976	56344	A	26117		2786	MAAVIEQIRRAVLALVTGVVHP GDIPVVWDIVWNATAAFIAVIII SLLLDESGFFEWAALHVLRWG YGPGRLPFNRIGLIRRITRRIRQR LLFGLVTFILLLNRSQNPAQDIQ LLAFRARTGKQTAQFIHHLPRM VFTDKTGSSNGLAPQGQCSAQ GELILNEKLAKQLVTAANWVK MQSDEGEINPVDILRWPGVMA AQEQDLDAIAAEILAALDGTLD DFIVARETEGQSGLKRVYHSPG APDIREFTRDAIP
25977 25978	56345	A	26118	48	219	EIFSVVWIIMTRGDVVTISRWR VSSSPLETWNKRWAKISPAPFS LLPISRLEKSEGDWLPETVISAF NMPSSLSLIISSL*FKVRDV*HFF /RI*TLRGHYRVINWPNFNIVVS QGIGKRLANSWLVFTLFHRI PETCGHLWAYVWPSCAAV\GL
237,0			20117			YFKVHVLG*RSVTPVTDIVKLL EFTRLRLPGYTKSIE
25979	56347	A	26120	1	912	

3498

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1	İ		sequence	1	
25980	56348	A	26121	1	1857	MRVCARACVRTRTRMCVYAH
23980	30348	A	20121	I	1037	TCVCVRTYAYVRVRVRVHVRA
ĺ		İ				RVRVCAYARTRVRNSLSILPFIQ
		į				LTLATPIHHIHQEEFNIRGIVPVL
			İ			RRVKPDLAIGIDITPSCDTPDLH
						DYSEVRINQGVGITCLNYHGRG
						TLAGLITPPRLIRMLEQTALEHN
		İ				IPVQREVAPGVITETGYIQLFLP
						GWEIGFSPLALLLAFLCSTSPGF
						GDPDGLGVIAYQDTVRPNAAT
						AISELNALAVKGVILTGDNPRA
						AAAIAGELGLEFKAGLLPEDKV
		İ				KAVTELNQHAPLAMVGDGIND
1		1				APAMKAAAIGIAMGSGTDVAL
						ETADAALTHNHLRGLVQMIEL
						ARATHANIRQNITIALGLKGIFL
						VTTLDDRVVAGSAGRYGGDGA
]				GDSECVKIVAQEIRQTDRNIER
						DRSPRPSGERVRVRGKGGIEAN
						QPLSTAFTNQITIRQSIRLFSNQF
			ı			VFHKGISRVVAGSSAIISNVSPG
		1				CILRTPNCRRGFPISCTGEGKAA
						YGWQRDGGHAEYLLAEEKDLI
ĺ	:					LLPDALSYEDGAFISCGVGTAY
						EGILRGEVSGSDNVLVVGLGPV
					11 13	GMMAMMLAKGRGAKRIIGVD
						MLPERLAMAKQLGVMDHGY*I
						LPDVY/V*IGVARVSWMNGRID
:						SELRTRVRAYAHTRTRARTCTR
						TRTRTYAYVRTHTHVCAYTHM
						RVRVRTHARAHTR
25981	56349	A	26122	3	1180	SKLGTRRSVVWA*SPSTSPTLW
23701	150517	'				CSTFSAAGHSSMKRMNEFVDL
	}					LPAQQRMKGENWYRGTADAV
ľ						TONLDIIRRYKAEYVVILAGDHI
						YKODYSRMLIDHVEKGARCTV
		-				ACMPVPIEEASAFGVMAVDEN
						DKIIEFVEKPANPPSMPNDPSKS
						LASMGIYVFDADYLYELLEEDD
						RDENSSHDFGKDLIPKITEAGLA
		1				YAHPFPLSCVQSDPDAEPYLAR
						CGYAGNFMESEPRSGLIRVSCM
						PAGPPLPLPAARSEAPKAAGTV
						ASVPSIAPARLRPVPDPVELVLV
						AAEFITPGDPTPRLHGSGFIDIR
					1	QIHHQTRSHLEGVKTGIRFLNH
]			}			FSGNPQGGIAHVNGVARFQVK
						QCHQAWGQQYAARLRFQARGI
		\perp				SLQIAIHRVDIIHRFDVRQL
25982	56350	Α	26123	16	543	
25983	56351	A	26124	1	3387	<u> </u>